Sequence 44, Appl Sequence 44, Appl Sequence 44, Appl Sequence 338, Appl Sequence 6, Appli Sequence 10, Appli Sequence 10, Appli Sequence 3, Appli Sequence 3, Appli Sequence 1, Appli Sequence 2, Appli Sequence 12, Appli Sequence 12, Appli Sequence 12, Appli Sequence 14, Appli Sequence 1, Appli

Sequence:

Run on:

Sequence 1 Sequence 1 Sequence 1

Sequence Sequence Sequence

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WS-09-300-958A-27

WS-09-300-958A-27

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 27, Application

Sequence 27, Application

Sequence 27, Application

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: Trenkle, Thomas

TITLE OF INVENTION: Using Same

TITLE OF INVENTION UNBER: US/09/300,958A

CURRENT APPLICATION NUMBER: US/09/301

PRIOR PAPLICATION NUMBER: 60/098,070

PRIOR PAPLICATION NUMBER: 60/118,624

PRIOR APPLICATION NUMBER: 60/118,624

PRIOR APPLICATION NUMBER: 80/118,624

NUMBER OF SEQ ID NOS: 85

SOFTWARE: PatentIn Ver. 2.0
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Mismatches:
Indels:
            US-09-352-616A-44
US-09-023-149A-44
US-09-009-913-8
US-09-009-913-6
US-09-009-913-6
US-09-016-434-927
US-08-86-281-1
US-08-878-177-3
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US-09-570-593-12
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US-09-344-579-1
US-09-360-779-1
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US-09-920-759-10
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Matches:
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92.72%
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Best Local Similarity:
                                                                                                                                           US-09-300-958A-27
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LENGTH: 1907
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Sequence 282, App.
Sequence 282, App.
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Sequence 4, Appli
Sequence 1, Appli
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Sequence 4, Appli
Sequence 44, Appli
                                                                              February 12, 2004, 19:34:21; Search time 93.2699 Seconds (without alignments) 1750.959 Million cell updates/sec
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Sequence 282, 3
Sequence 2, Api
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                     - nucleic search, using frame_plus_p2n model
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US-09-570-593-4
US-08-746-719A-1
US-09-389-681-282
US-09-339-318-282
US-09-333-326B-282
US-09-433-826B-282
US-09-620-405B-282
US-09-631-826B-282
US-09-601-287A-282
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US-09-009-913-4
US-09-020-956-44
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Listing first 45 summaries
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1980
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Result

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996 GGCGTCTTCAAGTTCCTGCGCTCCCAAGGCTGTGGCCCCAACTATGGGGCCAAAAGAAAAG 1055
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                                                           GluAspProThrLeuAlaProAlaProPro---ThrThrPheGlyThrGluAspLeuVal 39
                                                                                                                      LeuThrieuAsnAsnGlnGlnMetThrieuGluGlyProGluLysAlaSerTrpThrSer 59
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                                                                                159 GAGGACTCCACCCTGGCTCTGTTCCCCCTGCTGCCACTTTGGGGCCGATGACTTGGTA
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99 GCTGCAACCTGTGAGATTAGCAACATTTTTAGCAACTACTTCAGTGCGATGTACAGCTCG 158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Human epithelial-restricted with serine box (ESX) OTHER INFORMATION: protein.
                                                                                                             Sequence 4, Application US/09570593

Patent No. 656663

GENERAL INFORMATION:
APPLICANT: Raifmann, Joerg
APPLICANT: Raifmann, Joerg
APPLICANT: All, Honge
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
FILE REPERBNCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13

NUMBER OF SEQ ID NOS: 13

SEQ ID NO 4
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Matches:
Conservative:
Mismatches:
                        1176 TGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 1208
360 TrpLysGluGluGluValGlyGluSerArgAsn 370
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ORGANISM: Homo sapiens
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Best Local Similarity:
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NAME/KEY: CDS
LOCATION: (96).
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US-09-570-593-4
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160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyr 179
                       180 CysSerThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199
                                                                                                             636 CCCGGCAGCTGTGCCCCAGGAGCCCCCTCCCTGGCAGCTCTGACGTCTCCACCGCAGGG 695
                                                                                                                                                         200 ThrAlaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219
                                                                                                                                                                                696 ACTGGTGCTTCTCGGAGCTCCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTGGAT 755
                                                                                                                                                                                                                                           239
                                                                                                                                                                                                                                                              756 CCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGAGGGATCCC (815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       936 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 995
                                                                                                                                                                                                                                                                                                                                                                                                                              876 CTCGAGGGCAAGAAGAGCAAGCACCCAGAGGCACCACCTGTGGGAGTTCATCCGG
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                                                                                                                                                                                                                                                                                                                                                           816 AAGCACGGGAAGCGGAAACGAGGCCGGCCCGGAAGCTGAGCAAAGAGTACTGGGACTGT
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APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
COUNTRY: NA STATE: PA
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08746789A Patent No. 5789200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 180 CysSerThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                              Length:
                                       ATG 50024
            REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270 626
INFORMATION FELEPAS: 610 270 626
INFORMATION FOR SEC ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1920
TYPE: Nucleic Acid
STRANDEDNESS: Single
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1707.00
92.72%
86.79%
86.21%
William T. Han
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Best Local Similarity:
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| Sequence 282, Application US/09389681A |
| Sequence 282, Application US/09389681A |
| Sequence 282, Application US/09389681A |
| Patent No. 6518237 |
| GENERAL INFORMATION: Application: Davin C. |
| APPLICANT: Miccham, Jennifer L. |
| APPLICANT: Miccham, Jennifer L. |
| APPLICANT: Miccham, Jennifer L. |
| APPLICANT: W. Jiangchun |
| TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND METHODS FOR THEIR USE |
| TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE |
| TITLE OF INVENTION: UNMERER: US/09/389,681A |
| CURRENT FILING DATE: 1999-09-02 |
| NUMBER OF SEQ ID NOS: 463-09-02 |
| SOFTWARE: FastSEQ for Windows Version 3.0 |
| SEQ ID NO 282 |
| LENGTH: 502 |
                                                                                                                                                                                                                                                                               1015 GGCGTCTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCAACTATGGGGGCAAAAGAAAAG 1074
                                                                                                                                                                                                  1075 AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 1134
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240 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 259
                                                                                 260 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 279
                                                                                                                       895 CTCGAGGGGAAGAAGGCACGCGCCCAGAGGCACCCACTGTGGGAGTTCATCCGG 954
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Conservative:
Mismatches:
Indels:
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93.33$
90.30$
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; ORGANISM: Homo sapiens
US-09-389-681-282
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Best Local Similarity:
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Pred. No.:
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186 CGGAAACGAGGCCGCGGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGGCAAG 245
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) Sequence 282, Application US/09620405B
) Patent No. 6528054
) GENERAL INFORMATION:
| APPLICANT: Uiang, Yuqiu
| APPLICANT: Dillon, Davin C. |
| APPLICANT: Mitcham, Jennifer L. |
| APPLICANT: Mitcham, Jennifer L. |
| APPLICANT: Harlocker, Susan L. |
| APPLICANT: Harlocker, Susan L. |
| APPLICANT: Harlocker, William T. |
| TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE |
| TITLE OF INVENTION: COMPOSITIONS |
| FILE REPERENCE: 210121.470C8 |
| CURRENT APPLICATION NUMBER: US/09/620,405B |
| CURRENT APPLICATION NUMBER: US/09/620,405B |
| CURRENT APPLICATION NUMBER: US/09/620,405B |
| SOFTWARE: FastSEQ for Windows Version 3.0 |
| SEQ ID NO 282 |
| LINGTH: 502
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ORGANISM: Homo sapiens
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304 PheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysAsnSerAsnMet 323
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Mismatches:
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Sequence 282, Application US/09433826B

; Patent No. 6579973

; GENERAL INFORMATION:
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Best Local Similarity:
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264 LysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArgAsplleLeulle 283
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APPLICANT: Vigiu, Jiang
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Xu, Jiangchun ComPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
CURRENT APPLICATION WUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FASTSEQ for Windows Version 3.0
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ORGANISM: Homo sapiens
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LENGTH: 502
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306 CACCCGGAGCTCAACGAGGCCTCATGAAGTGGGGAAGGCATGAAGGCGTCTTCAAG 365
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APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C4
CURRENT FILING DATE: 1999-11-03
CURRENT FILING DATE: 1999-11-03
SOFTWARE: FastSEQ for Windows Version 3.0
SOFTWARE: FastSEQ for Windows Version 3.0
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324 ThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGluIleLeuGluArg 343
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Matches:
Conservative:
Mismatches:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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APPLICATION NUMBER: US/09/009,913
FILING DATE: 21-JAN-1998
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                                                                                                                                                                                                                                                                                                                        ; Sequence 2, Application US/09009913; Patent No. 6087485; GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                      344 ValAspGlyArgArg 348
                                                                                                                                                                                                  486 Grechrececedeca 500
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SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
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ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 650-327-3231
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US-09-009-913-2
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                                                                                                                                                                                                                                                            RESULT 9
US-09-009-913-2
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324 ThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlulleLeuGluArg 343
                                                                                                                                                426 ACCTACGAGAAGCTGAGCCGATGAGGTACTACTACAAACGGGAGATCCTGGAACGG 485
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    6 GGCGCAGGAGCCCCCCCCCCGGCAGCTCTGACGTCTCCACCGCAGGGAACTGGCTTCT

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APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: A. Jiangohun
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND FITLE REPERENCE: 210121.470C7
CURRENT APPLICATION HUBBER: US/09/604,287A
CURRENT FILING DATE: 2000-06-22
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Conservative:
Mismatches:
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 282, Application US/09604287A Patent No. 6586572 GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-604-287A-282
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426 ACCTACGAGAGAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGAACGG 485
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GENERAL INFORMATION:

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                                                                                 375 CAAGAGTTGGACATCAACGGCGAGCACCTCTGCAGCATGAGTTTGCAGGAGTTCACCCGG 434
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Sequence 3, Application US/09009913 Patent No. 6087485

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Mismatches:
                                                                                                                                                                     ZIP: 94301
COMPUTER READALE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
FILING DATE: 21-JAN-1998
CLASSIFICATION:
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Matches:
APPLICANT: Axys Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
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                                                                           ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
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SEQUENCE CHARACTERISTICS:
LENGTH: 5510 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,
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54.43%
40.82%
29.29%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: cDNA
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Best Local Similarity:
                                                                                                                                                         USA
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                                                                                                                                    STATE: C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AlaValAlaGlnLeuTrpGlyGlnLysLysAsnSerAsnMetThrTyrGluLysLeu 328
 174 GlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerProGlySerSer 193
                                                                        194 AspValSerThrAlaArgThrAlaThrProGlnSerSerHisAlaSerAspSerGlyGly 213
                                                                                                             -----ThrAspTyrLysLysGlyGluProLysHisGlyLysArgLysArgGlyArg 248
                                                                                                                                                                                                                                                                                                                                                                                          704 ------GIAGAITIGITGGACAGCAAAACITICIGCGGGCICAGAICTICGIG 751
                                                                                                                                                                                                                                                         752 ACAACCACCAGTCACCTTCCTGTTGCAGAGTCACCTGATATGAAAAGGAGCAAGACCC 811
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             962 GCAGTGGCTCAGCTATGGGGTAAAAAGAAGAAGAACAACAGCAGCATGACCTATGAAAAGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     349 LeuvalTyrLysPheGlyLysAsnSerSerGlyTrpLysGluGluGlu 364
                           665 GACGAGAACTATTATATGACACCAACTATGGT-------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDLIM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASISEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
FILING DATE: 21-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/09009913
Patent No. 6087485
GENERAL INFORMATION:
APPLICANT: AXYS PARTMECULICALS:
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sherwood, Pamela J
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 285 Ham:
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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US-09-009-913-4
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1059 CCAGGATTAATAAAATGGGAAGCACCGATCTGAGGGCGTCTTCAGGATTCTTGAAATCAGAG 1118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              134 LeuSerTrpllelleGluLeuGluLysAspGlyMetSerPheGlnGluSerLeuGly 153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   174 GlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerProGlySerSer 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  855 -----AGCACA----- 860
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    909 ACAACCACCAGTCACCTTCCTGTTGCAGAGTCACCTGATATGAAAAAGGAGCAAGACCCC 968
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Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                              Length:
                                                                                                                                                                                                                                                                                                                                                                                    Gaps:
                                                          INFORMATION FOR SEQ ID NO: 4: SEQUENCE WHARACTERISTICS: LENGTH: 5667 base pairs TYPE: nucleic acid STRANDENESS: double TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                            2.17e-51
                                                                                                                                                                                                                                                                                              580.00
54.43%
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                    TELEFAX: 650-327-3231
                                                                                                                                                                                                                                                                                                                     Percent Similarity:
Best Local Similarity:
TELEPHONE:
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GREERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
     109 ArgLeuValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThrSer 128
                                           730 ACCCGGGCGGCGGGGAGCANCTCCTCTACAGCAACTTGCAGCATCTGAAGTGG 671
                                                                                                                                                                                                   146 MetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAla 165
                                                                                                                                                                                                                                                                                                 166 GlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyPro 185
                                                                                                                                                                                                                                                                                                                                                                                            186 GlyAlaProSerProGlySerSerAspValSerThrAlaArgThrAlaThrProGlnSer 205
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                                                                                                                                                     670 AACGGCCAGTGCAGTAGTGAC
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APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR INMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                1119 GCAGTGGCTCAGCTATGGGGTAAAAGAAGAACAACAGCAGCATGACCTATGAAAAGCTC 1178
                                                                                                                                           SerArgAlaMetArgTyrTyrTyrLysArgGluIleLeuGluArgValAgDGlyArgArg 348
309 AlavalAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeu 328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NDATA:
APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
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Indels:
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Patent No. 6261562
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NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFRENCE/DOCKET NUMBER: 21012
REFRENCE/DOCKET NUMBER: 21012
TELECOMMUNICATION INFORMATION:
TELEFAX: (206) 622-4900
TELEFAX: (206) 622-6911
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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ZIP: 98104
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NAME/KEY: misc feature
LOCATION: (1)...(852)
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ORGANISM: Homo sapien
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Query Match:
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US-09-439-313-44/c
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                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/030,607 FILING DATE: 25-FEB-1998 CLASSIFICATION:
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Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length:
Matches:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: Maki, David J.
REGIETRATION NUMBER: 31,392
REFRENCE/DOCKET NUMBER: 21012
TELECOMMUNICATION INFORMATION:
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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54.28%
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STRANDEDNESS: single
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Best Local Similarity:
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MOLECULE TYPE: CL
ORIGINAL SOURCE:
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364 ATCCTCTIGAACCCAGACAAGAACCCAGGATTAATAAAATGGGAAGACCGATCTGAGGGC 305
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241 HisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCysLeu 260
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                                                                                                                                                       409 -------AAGCACAACCCGAQAGGACTCACTTATGGAATTCATCCGCGAC
                                     261 GluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArgAsp
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APPLICANT: Read, Steven G.
APPLICANT: Railor, Michael
APPLICANT: Renger, Gary
APPLICANT: Roll, John
APPLICANT: Bolk, John
APPLICANT: Day, Craig
TILLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TILLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210.12.42.769
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT PILLING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFFWARE: RestSEQ for Windows Version 3.0
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Matches:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
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IHER INFORMATION: n = A,T,C or G
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Conservative:
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NAME/KEY: misc_feature
LOCATION: (1)...(852)
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ORGANISM: Homo sapien
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146 MetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAla 165
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Sequence 44, Application US/09352616A

Sequence 44, Application US/09352616A

Sequence 44, Application US/09352616A

GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: W, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
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TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT PAPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
SEQ ID NO 44
LENGTH: 852
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rGluAlaValAlaGlnLeuTrpGly	304 GTCTCAGGTTCTTGAAATCAGAGGCAGTGGCTCAGCTATGGGGTAAAAGAAGAACAAC 245	321 SerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlulle 340	244 AGCAGCATGACCTATGAAAAGCTCAGCGAGCTATGAGATATTACTACAAAAGAGAAATT 185	341 LeuGluargValaspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGlyTrp 360		361 Lysglugluglu 364	124 AGAGAAATGAA 113	Search completed: Pebruary 12, 2004, 21:50:15 Job time : 105.27 secs
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Patent No. 6495319

GENERAL INPORMATION:
APPLICANT: Welsh, John

APPLICANT: Welsh, John

APPLICANT: Trenkle, Thomas

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE REPRESENCE: P-PH 3457

CURRENT FILING DATE: 1999-04-27

PRIOR APPLICATION NUMBER: 60/098,070

PRIOR PILING DATE: 1998-08-27

PRIOR PILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-02-04

NUMBER OF SEQ ID NOS: 85

SEOFWARE: PATENTIN VEL 2.0
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Sequence 182, App
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66.7%; Pred. No. 2.3e-49;
iive 0; Mismatches 195; Indels 33; Gaps
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US-09-302-769-1
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US-09-449-218D-17
US-09-668-529A-17
US-09-668-529A-17
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US-09-668-529A-17
US-09-668-529A-17
US-09-68-529A-17
US-09-68-529A-17
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US-09-68-529A-17
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Best Local Similarity 66.7°
Matches 456; Conservative
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ORGANISM: Homo sapiens
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US-09-300-958A-27
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7239 GAAGAACCIGITGGGCGIGCCCICTGC---AGICTCCTCAAGIGCAGCCITTGGCCIC--

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1394 CICCCCACCCTCCTCTGGAATTACAAGCCCTGGGGTTTGAAGCTGACTTTATAGCTGCA 1453
                                                                                                                                                      1334 AAGCTGATGTTTTGGTGTATTGTCAGCCATCGTCCTTGGACTCGAGACTATGGCCTCGC 1393
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                                                       7457 AGACTGCAGGGACGGAGGGACAGGGTTGTGTCCTCGGTACTTC---CTGGACTGCCTTC 7513
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Sequence 1, Application US/08746789A
Setent No. 5789200
GENERAL INFORMATION:
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
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APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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TELEPHONE: 610 270 5219
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INFORMATION FOR SEQ ID NO: 1:
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ATTORNEY/AGENT INFORMATION:
NAME: William T. Han
REGISTRATION NUMBER: 34,3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        King of Prussia
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                                                                                                                                                                                                                                          1514 AACACCTTCTTCCTGCAGACACTTGGACTGAGCCAAGGAGGCTTGGGAGGCCT----AG 1569
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                                            1394 CTCCCCACCCTCCTCTTGGAATTACAAGCCCTGGGGTTTGAAGCTGACTTATAGCTGCA 1453
                                                                                                                                              1454 AGTGTATCTCCTTTTATCTGGTGCCTCCTCAAACCCAGTCTCAGACACTTAAATGCAGAC 1513
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66.7%; Pred. No. 2.3e-49;
ative 0; Mismatches 195; Indels 33; Gaps
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; OTHER INFORMATION: Human epithelial-restricted with serine box (BSX)
; OTHER INFORMATION: protein.
US-09-570-593-4
7294 --TCTCCTCGCCCTCTTGGAATTACAAGCCCCGGGTTTGAACCAACTTGT-----
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APPLICANT: Kanfmann, Joerg
APPLICANT: Kin, Hong
APPLICANT: Harrowe, Greg
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOWAIN PROTEINS IN
FILE REFERENCE: 2300-1556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. 6566063
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Best Local Similarity
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6999 AGGTATTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7294 --TCTCCTCGCCCTCTTGGAATTACAAGCCCCGGGTTTGAACCAACTTGTTCGA---- 7345
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PREACH NO. 6518237
GENERAL INFORMATION:
APPLICANT: Yuqui, Jiang
APPLICANT: Micham, Davin C.
APPLICANT: Micham, Dennifer L.
APPLICANT: Xu, Jiangchun
TILLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TILLE OF INVENTION: DAMPOSITIONS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C3
                                                                                                                                                                                                                   Query Match 2.8%; Score 215.6; DB 1; Length 1920; Best.Local Similarity 65.6%; Pred. No. 3e-49; Matches 449; Conservative 0; Mismatches 204; Indels 31;
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                       LENGTH: 1920

TYPE: Nucleic Acid

STRANDEDNESS: Single

TOPOLOGY: Linear

ANTI-SENSE: NO
SEQUENCE CHARACTERISTICS
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6256 GCCCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGGAGCTC 6315
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Best Local Similarity 88.9%; Pred. No. 2.9e-35;
Matches 177; Conservative 0; Mismatches 22; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
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APPLICANT: Jillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitchen, Jennifer L.
TITLE OF INVENTION: DIAGNOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSITIONS OF BREAST CANCER
FILE REFERENCE: 21011.47008
CURRENT APPLICATION NUMBER: US/09/620,405B
CURRENT APPLICATION NUMBER: 2000-07-20
NUMBER OF SEQ ID NOS: 495
SOFTWARE: FREESEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                Query Match 2.1%; Score 163.8; DB 4; Length Best Local Similarity 88.9%; Pred. No. 2.9e-35; Matches 177; Conservative 0; Mismatches 22; Indels
CURRENT APPLICATION NUMBER: US/09/389, 681A CURRENT FILING DATE: 1999-09-02 NUMBER OF SEQ ID NOS: 463 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ) Sequence 282, Application US/09620405B ; Patent No. 6528054
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                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
US-09-389-681-282
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CRGANISM: Homo sapiens
US-09-620-4058-282
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LENGTH: 502
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; ORGANISM: Homo sapiens
US-09-604-287A-282
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Patent No. 6579973

GENERAL INFORMATION:
APPLICANT: Jiang, Yuqui
APPLICANT: Dillon, Davin C.
APPLICANT: Michan, Jennifer L.
APPLICANT: Wi Jiangchun COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT APPLICATION NUMBER: 1999-11-03
NUMBER OF SEQ ID NOS: 474
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                                                                                                                                                                                          Sequence 282, Application US/09339338A

Batent No. 6573368

General information of the partial o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         n 2.1%; Score 163.8; DB 4; Length 502; Similarity 88.9%; Pred. No. 2.9e-35; 77; Conservative 0; Mismatches 22; Indels 0
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438 CIGAGCCGGGCCATGAGGT 456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-339-338-282
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US-09-433-826B-282
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                                                                                                                   RESULT 6
US-09-339-338-282
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6376 GAGGCCGTGGCCCAACTCTGGGGCCAGAAGAAGAAGAACAGCAACATGACCTATGAGAAG 6435
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                                                                                                                                              6316 AACGAAGGCCTCATGAAGTGGGAGAACCGGCACGAGGGTGTGTTCAAGTTTCTTCGCTCA 6375
                                                                                                                                                                                                                                                                                            6376 GAGGCCGTGGCCCAACTCTGGGGCCAGAAGAAGAAGAACAGCAACATGACTTATGAGAAG 6435
6256 GCCCCCAGAGGIACICACCIGIGGGAGIIITAICCGAGACAICCIAAICCACCCCGAGGIC 6315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             318 AACGAGGGCTCATGAAGTGGGAAATCGGCATGAAGGCGTCTTCAAGTTCCTGCGCTCC 377
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                                                                                                                                                                                                               318 AACGAGGGCCTCATGAAGTGGGAAATCGGCATGAAGGCGTCTTCAAGTTCCTGCGCTCC 377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Mitcham, Jannifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Ku, Jiangchun
APPLICANT: Ku, Jiangchun
APPLICANT: Ku, Jiangchun
APPLICANT: Ku, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.47007
CURRENT FILLING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FASTSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING TITLE OF INVENTION: PATHWAY GENE EXPRESSION NUMBER OF SEQUENCES: 1450
CORRESPONDENT ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.1%; Score 163.8; DB 4; Length 502; Best Local Similarity 88.9%; Pred. No. 2.9e-35; Matches 177; Conservative 0; Mismatches 22; Indels 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 282, Application US/09604287A Patent No. 6586572
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Patent No. 6500938
GENEAL INFORMATION:
APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6436 CTGAGCCGAGCCATGAGGT 6454
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Sequence 4, Application US/09245041

Sequence 4, Application US/09245041

Patent No. 6274339

GENERAL INFORMATION:
APPLICANT: Magle, D.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT
TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
FILE REPERENCE: 7883-136
CURRENT APPLICATION NUMBER: US/09/245,041

CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/093,630
EARLIER FILING DATE: 1998-07-21
EARLIER APPLICATION NUMBER: 60/104,978
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                                                                                                                                                                                                                                        SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
SSEE: INCYTE PHARMACEUTICALS, INC.
I: 3174 PORTER DRIVE
PALO ALTO
CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0002 US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1050 845-4166
INFORMATION FOR SEQ ID NO: 927;
SEQUENCE CHARACTERISTICS:
                                                                                                                 ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEB PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 79.3
Matches 149; Conservative
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CLONE: 773734
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       ADDRESSEE:
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                                                                                                     COUNTRY:
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US-09-020-956-44/c
US-09-020-956-44/c

J Sequence 44, Application US/09020956

J Patent No. 6261560

GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Nauher OF SEQUENCES: 178

CORRESPONDENCE ADDRESS:
ADDRESSER: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                           5390 ITCTITITICATITICGGTTTTTCAAGACAGGGTTTCTCTGTATAGCCCTGGCTGTCCTG 5449
                                                                                                                                                                                                                                                                                                                                                      1841 TTTGTTCTGTAGTTTTTAGTTTTTCAAGACAGAGTTTCTCTGTGTAGCCCTGGCTGTCCTG 1900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1961 GAGTGCTGGGATTAAAGATGTGCGCTGCCCTCCTCCACCCCAATTGTTTTTGTTTTTTA
                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                            Length 5973;
                                                                                                                                                                                                                                      Query Match
1.6%; Score 123.6; DB 3; Length Best Local Similarity 67.4%; Pred. No. 2e-23;
Matches 174; Conservative 0; Mismatches 84; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC Compatible
OPERATUNG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Mark, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 812 base pairs
                         NUMBER OF SEQ ID NOS: 131
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2081 TGTTGGGCTTCTGACCAC 2098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5630 CATTTTTTTCATTCTAC 5647
EARLIER FILING DATE: 1998-10-20
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TYPE: nucleic acid
STRANDEDNESS: single
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                 TYPE: DNA ORGANISM: Mus musculus
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                                                                          SEQ ID NO 4
LENGTH: 5973
                                                                                                                                                                                     US-09-245-041-4
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6378 GGCCGTGGCCCAACTCTGGGGCCAGAAGAAGAAGAACAGCAACATGACCTATGAGAAGCT 6437
                                                                     6318 CGAAGGCCTCATGAAGTGGGAAAACCGGCAAGGGTGTGTTCAAGTTTCTTCGCTCAGA 6377
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                                                                                                                                                                                                 281 GGCAGTGAGCTCAGCTATGAGAAAGAACAACAACAACAGCATGACCTATGAAAAGCT 222
                       401 cccgagagagacrcacrrargagaarrcarccacaacarccrcrrgaacccaagaa 342
                                                                                                                    341 cccaddarraaraaraddaaqaccdarcrdadddcdrcrrcaddrrcrrgaaarcada 282
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APPLICANT: Read, Stewen G.

APPLICANT: Raides, Michael
APPLICANT: Refed, Michael
APPLICANT: Refer, Mark
APPLICANT: Refer, Mark
APPLICANT: Solk, John
APPLICANT: Solk, John
APPLICANT: OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFREENCE: 210121.42769
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FREISEQ FOR WINDOWS Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 852;
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74.1%; Pred. No. 1.1e-21;
tive 0; Mismatches 51;
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US-09-352-616A-44/c
; Sequence 44, Application US/09352616A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: ALtcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang Yuqui
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 44, Application US/09439313; Patent No. 6329505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (1)...(852)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                  6438 GAGCCGAGCCATGAGGT 6454
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                                                                                                                                                                                                                                                                                                             221 CAGCCGAGCTATGAGAT 205
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Best Local Similarity 74.14
Matches 146; Conservative
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Sequence 44, Application US/09030607
Patent No. 626245
GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
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                                                                                                                                                                                            6258 CCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGAGCTCAA 6317
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                                                                                                                                             0; Gaps
                                                                                            Query Match 1.5%; Score 115.4; DB 3; Length 852; Best Local Similarity 74.1%; Pred. No. 1.1e-21; Matches 146; Conservative 0; Mismatches 51; Indels 0
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1.5%; Score 115.4; DB 3; Length 852;
Best Local Similarity 74.1%; Pred. No. 1.1e-21;
Matches 146; Conservative 0; Mismatches 51; Indels 0
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: RatentIn Release #1.0, Version #1.30
APPLICATION NUMBER: US/09/^^*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 2101.
TELECOMUNICATION INDORMATION:
TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
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CLASSIFICATION:
                       ) ORGANISM: Homo sapiens
US-09-020-956-44
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ORIGINAL SOURCE:

ORGANISM: Homo sag

US-09-030-607-44
                                                                                          Query Match
Best Local Similarity
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ORIGINAL SOURCE:
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us-08-978-217-15.rni

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6258 CCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGAGCTCAA 6317
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GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Micham, Jennifer Lynn
APPLICANT: Micham, Jennifer Lynn
APPLICANT: Micham, Jennifer Lynn
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
TILE REPERENCE: 210121.427C8
CURRENT PAPLICANTON NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 1.5%; Score 115.4; DB 4; Length 852; Best Local Similarity 74.1%; Pred. No. 1.1e-21; Matches 146; Conservative 0; Mismatches 51; Indels 0
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US-09-232-149A-44/C

Sequence 44, Application US/09232149A

Sequence 44, Application US/09232149A

Setent No. 6465611

GENERAL INFORMATION:

APPLICANT: W. Jiangchun

APPLICANT: Dillon, Davin C.

APPLICANT: Mitcham, Jennifer Lynn

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE

TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE

FILE REPERENCE: 210121.427C6

CURRENT APPLICATION NUMBER: US/09/232,149A

CURRENT FILING DATE: 1999-01-15

NUMBER OF SEQ ID NOS: 338

SOFTWARER FEASTSEQ for Windows Version 3.0
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Best Local Similarity 74.1%; Pred. No. 1.1e-21;
Matches 146; Conservative 0; Mismatches 51; Indels 0;
                                                                                                                                                                                                                                                                                                                       FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LCCATION: (1)...(852)
CTHER INFORMATION: n = A,T,C or G
US-09-352-616A-44
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LCCATION: (1)...(852)

/ OTHER INFORMATION: n = A,T,C or G

US-09-232-149A-44
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SEQ ID NO 44
LENGTH: 852
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LENGTH: 852
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5258 CCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGAGCTCAA 6317
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                             401 CCCGAGAGACTCACTTATGGGAATTCATCCGCGACATCCTCTTGAACCCAGACAAGAA 342
                                                                                                                                                              341 CCCAGGATTAATAAAATGGGAAGACCGATCTGAGGGCGTCTTCAGGTTCTTGAAATCAGA 282
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Sequence 17, Appli
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Sequence 384, App
Sequence 1, Appli
Sequence 55, Appl
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Sequence 338, App
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7673.534 Million cell updates/sec
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                                                                                                                                                                                           February 12, 2004, 21:46:16 ; Search time 1.20792 Seconds
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-389-681-282
US-09-620-405B-282
US-09-433-826B-282
US-09-604-287A-282
US-09-570-593-4
US-09-570-593-4
US-08-746-789A-1
US-08-746-789A-1
US-08-746-789A-1
US-08-746-789A-1
US-08-479-72B-1
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US-08-479-72B-1
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3 US-09-103-840A-1

US-09-123-840A-1

US-09-123-030-29

US-09-522-17-88

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US-09-521-017B-384

US-09-519-353-1
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US-09-495-050A-190
US-09-009-913-338
US-09-535-008-55
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                            OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Patent No. 6500338

GENERAL INFORMATION:

APPLICANT: Janica Au-Young

APPLICANT: Janica Au-Young

APPLICANT: Janica Au-Young

TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING

TITLE OF INVENTION: PATHWAY GENE EXPRESSION

NUMBER OF SEQUENCES: 1490

CORRESPONDENCE ADDRESS:

ADDRESSEE: INCYTE PHARMACEUTICALS, INC.

STREET: 3174 PORTER DRIVE

STREET: CALIFORNIA

COUNTRY: USA
                                     Sequence
Sequence
Sequence
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US-09-708-200-16
US-08-468-024B-1
US-08-187-757D-1
US-09-620-312D-240
US-09-620-312D-240
US-09-333-599-5
US-09-333-599-5
US-09-499-781-5
US-09-499-781-1
US-09-499-781-1
US-09-499-781-1
US-09-499-781-1
US-09-499-781-1
US-09-499-781-1
US-09-499-781-1
US-09-99-781-1
US-09-99-781-1
US-09-99-781-1
US-09-99-781-1
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION WIMBER:
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION WIMBER: 37 071
REFERENCE/DOCKET NUMBER: PA-0002 US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION ON SEQ ID NO: 927:
SEQUENCE CHARACTERISTICS:
LENGTH: 237 Dase pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: STRANDEDNESS: single
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Pred. No. (
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Best Local Similarity
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; CLONE: 773734
US-09-016-434-927
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US-09-433-826B-282
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LENGTH: 502
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Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

APPLICANT: Vuqui, Jiang

APPLICANT: Vuqui, Jiang

APPLICANT: Xi, Jiangchun

TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND

TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE

FILE REFERENCE: 210121.470C3

CURRENT APPLICATION WINHER: US/09/389,681A

CURRENT FILING DATE: 1999-09-02

NUMBER OF SEQ ID NOS: 463

SOFTWARE: FREESE FREESE FOR Windows Version 3.0
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Sequence 282, Application US/09620405B

Patent No. 628054

GENERAL INTORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: William L.
APPLICANT: Harlocker, Susan L.
APPLICANT: William L.
TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
FILE REPERENCE: 210121.47068

CURRENT APPLICATION NUMBER: US/09/620,405B

CURRENT FILING DATE: 2000-07-20

NUMBER OF SEQ ID NOS: 495

SOFFWARDER: Pastern of Containing Con
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100.0%; Score 21; DB 4; Length 502;
Best Local Similarity 100.0%; Pred. No. 0.31;
Matches 21; Conservative 0; Mismatches 0; Indels
    0; Mismatches
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LENGTH: 502
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21; Conservative
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CRGANISM: Homo sapiens
US-09-389-681-282
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ORGANISM: Homo sapiens
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US-09-339-338-282/c

RESULT 4

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APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Juqui
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangcham, Jennifer L.
APPLICANT: Xu, Jiangcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121-470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT APPLICATION NUMBER: US/09/433,826B
SUPTABER FILING DATE: 1999-11-03
SUPTABER FILING DATE: 1999-11-03
SUPTABER: FastSEQ for Windows Version 3.0
Sequence 282, Application US/09339338A
Patent No. 657356
Patent No. 657356
Patent No. 657356
APPLICANT: Virgiu, Jiang
APPLICANT: Wigiu, Jiang
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: AL, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE FREE FREESE FREESE FOR Windows Version 3.0
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APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 21; DB 4; Length 502; 100.0%; Pred. No. 0.31;
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; Sequence 282, Application US/09433826B
; Patent No. 6579973
; GRNERAL INFORMATION:
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; Sequence 282, Application US/09604287A
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GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.0
Matches 21; Conservative
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US-09-339-282
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SOFTWARE: FastSEQ for Windows Version 4.0

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US-09-300-958A-27/c

| Sequence 27, Application US/09300958A |
| Patent No. 6495319 |
| Patent No. 6495319 |
| GENERAL INFORMATION |
| APPLICANT | Welsh, John | |
| TITLE OF INVENTION | Reduced Complexity Nucleic Acid Targets and Methods of TITLE OF INVENTION | Using Same |
| FILE REPERENCE | P-PH 3457 |
| CURRENT FILING DATE: 1999-04-27 |
| PRIOR FILING DATE: 1998-04-27 |
| PRIOR FILING DATE: 1998-03-27 |
| PRIOR FILING DATE: 1998-03-27 |
| PRIOR FILING DATE: 1998-02-27 |
| PRIOR FILING DATE: 1998-02-04 |
| NUMBER OF SEQ ID NOS: 85 |
| SOFTWARE | PATENTIN OFF 2.0
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APPLICANT: Kaufmann, Joerg
APPLICANT: Kin, Hong
APPLICANT: Atin, Hong
APPLICANT: Harrowe, Greg
ITILE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
ITILE OF INVENTION: CANCER
FILE REPERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER FILE REFERENCE: 210121.470C7
CURRENT PELLING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 22
LENGTH: 502
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Patent No. 6566063
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Matches 21, Conservative
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Best Local Similarity 100.
Matches 21, Conservative
                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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LENGTH: 1907
                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                               TYPE: DNA
ORGANISM: Homo sapiens
ORGANISM: Homo sapiens
INAME/KEY:
NAME/KEY:
OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
US-09-570-593-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08746789A
Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
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1907;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 21, DB 1, Length 1920; 100.0%; Pred. No. 0.33; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IMM 486
COMPUTER: IMM 280
COMPU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1100 Graccrcardeccccddcrcag 1080
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ATTORNEY/AGENT INFORMATION:
NAME: William T. Han
REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELERHONE: 610 270 5219
TELEFAX: 610 270 4026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      34,344
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Matches 21; Conservative
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LENGTH: 1920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: Linear ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JS-08-746-789A-1/c
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SEO ID NO 4
LENGTH: 1907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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us-08-978-217-14.rni

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) NAME/KEY: CDS
) LOCATION: (316)..(1602)
US-09-213-767-1
                 STREET: F.C.
                                                                      Texas
                                                                                COUNTRY: Un
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION:
PCT-US95-02251-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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| Sequence 17, Application PC/TUS9502251
| GENERAL INFORMATION: GENERAL INFORMATION: TITLE OF INVENTION: TITLE OF INVENTION: CELLS
| TITLE OF INVENTION: CELLS
| NUMBER OF SEQUENCES: 18
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 78.1%; Score 16.4; DB 3; Length 5499; Best Local Similarity 94.4%; Pred. No. 49; Matches 17; Conservative 0; Mismatches 1; Indels 0.
                                                                                  GENERAL INFORMATION:
APPLICANT: Bonadio, Jeffrey
APPLICANT: Win, Wushan
TITLE OF INVENTION: LATENT TGF( BINDING PROTEIN (LTBP)
TITLE OF INVENTION: GENES, COMPOSITIONS AND METHODS
NUMBER OF SECUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TEB PC compatible
COMPUTER: TEB PC compatible
COMPUTER: TEB PC compatible
COMPUTER: TEB PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURSTRING SYSTEM: PC-DOS/MS-DOS
CURSTRING MUMBER: US/08/479,722B
FILING DATE: 07-JUN-1995
CLASSIFICATION:
PRIOR APPLICATION NUMBER: US PCT/US95/02251
FILING DATE: 21-FEB-1995
FILING DATE: 30-SEP-1994
FILING DATE: 30-SEP-1994
FILING DATE: 18-FEB-1994
APPLICATION NUMBER: US 08/199,780
FILING DATE: 18-FEB-1994
ATTORNEY AGENT INFORMATION:
NAMME: FUSSEY, Shelley P.M.
REGISTRATION NUMBER: 31-458
REFERENCE/DOCKET NUMBER: 4100.000500/FUS
TELLECOMMUNICATION NUMBER: 31-458
REFERENCE/DOCKET NUMBER: 4100.000500/FUS
TELLECOMMUNICATION NUMBER: 31-458
                                                                                                                                                                                                                                              ADDRESSEE: Williams, Morgan & Amerson
STREET: 7676 Hillmont, Suite 250
RESULT 10
US-08-479-722B-1/c
Sequence 1, Application US/08479722B
Patent No. 6074840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3158 ACCTCATAGCCCGGCTCA 3141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (713) 934-7000
TELEPAX: (713) 934-7011
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 5499 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 5499 base pai
TYPE: nucleic acid
STRANDEDNESS: single
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; LOCATION: 1..5499
US-08-479-722B-1
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                                                                                                                                                                                                                                                                    STREET: 7676 H
                                                                                                                                                                                                                                                                                            STATE: Tex...
COUNTRY: USA
77040
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US-09-213-767-1/C

Sequence 1, Application US/09213767

Sequence 1, Application US/09213767

Sequence 1, Application US/09213767

GENERAL INFORMATION:

APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: ANTISENSE MODULATION OF ELK-1 EXPRESSION

TITLE OF INVENTION: ANTISENSE WOULATION OF ELK-1

CURRENT FILING DATE: 1998-12-17

SUMBER OF SEQ ID NOS: 47

LENGTH: 2266
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                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPBEATING SYSTEM: PC-DOS/MS-DOS/ASCII
SOFTWARE: Parentin Release #1.0, Version
SOFTWARE: #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/02251
FILING DATE: CONCURRENTLY HEREWITH
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,650
FILING DATE: 30-SEP-1994
CLASSIFICATION NUMBER: US 08/199,780
FILING DATE: 18-FEB-1994
CLASSIFICATION NUMBER: 32,165
REGISTATION NUMBER: 32,165
REGISTATION NUMBER: UNICO09P--
TELECOMMUNICATION INFORMATION:
ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
                                                                                                                     United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3158 ACCTCATAGCCCGGCTCA 3141
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INPORMATION POR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 5502 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 94.44
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: 1..5502
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Search completed: February 13, 2004, 01:55:01 Job time : 20.2079 secs
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US-09-103-840A-1
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OTHER INFORMATION: "n" bases at various positions throughout the sequence
OTHER INFORMATION: represent a, t, c or g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTHER INFORMATION: CDC 1551
GOTHER INFORMATION: "n" bases at various positions throughout the sequence of CTHER INFORMATION: represent a, t, c or g
US-09-103-840A-2
                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: FIELSCHMAN, Robert D.
APPLICANT: FIELSCHMAN, Robert D.
APPLICANT: FIELSCHMAN, Robert D.
APPLICANT: FRASER, Owen R.
APPLICANT: FRASER, John C.
APPLICANT: VENTER, John C.
APPLICANT: VENTER, John C.
APPLICANT: VENTER, John C.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: TUBRECULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, Oldn C.
TITLE OF INVENTION: DAG SCHENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
TITLE OF INVENTION: TUBERCULOSIS
FILE REPERENCE: 24366-22007.00
CURRENT APPLICATION UNDER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN Ver. 2.1
LENGTH: 4403765
                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 77.1%; Score 16.2; DB 3; Length 4403765; Best Local Similarity 85.7%; Pred. No. 54; Matches 18; Conservative 0; Mismatches 3; Indels 0;
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0
Query Match 77.1%; Score 16.2; DB 2; Length 2266; Best Local Similarity 85.7%; Pred. No. 59; Matches 18; Conservative 0; Mismatches 3; Indels 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Mycobacterium tuberculosis
                                                                                                                                                   513 GTACCGCAAGGCCCCGGCTGAG 493
                                                                                                  1 GTACCTCATGGCCCGGCTCAG 21
                                                                                                                                                                                                                                                                                Sequence 2, Application US/09103840A Patent No. 6294328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09103840A Patent No. 6294328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GTACCTCATGGCCCGGCTCAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: FLEISCHMAN, Robert D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77.1%;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 2
LENGTH: 4403765
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US-09-103-840A-2/c
                                                                                                                                                                                                                               RESULT 13
US-09-103-840A-2
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APPLICANT: FRASER, Claire M.
APPLICANT: FRASER, Claire M.
APPLICANT: FRANCH.
TITLE OF INVENTION: DAM SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-220007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN Ver. 2.1
  0; Gaps
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85.7%; Pred. No. 54;
tive 0; Mismatches 3; Indels 0;
3; Indels
  0; Mismatches
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ORGANISM: Mycobacterium tuberculosis
OTHER INFORMATION: H37Rv
US-09-103-840A-1
                                                                                                                                                                                                                           Sequence 1, Application US/09103840A
Patent No. 6294328
GENERAL INFORMATION:
APPLICANT: FLEISCHWAN, Robert D.
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Best Local Similarity 85.7%
Matches 18, Conservative
  Matches 18; Conservative
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Sequence 282, Application US/09389681A Patent No. 6518237 GENERAL INFORMATION:
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                                                                                                  , ORGANISM: Homo sapiens
US-09-389-681-282
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Matches 21; Conserv
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LENGIH: 502
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Sequence 282, App
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                                                                                                                                 February 12, 2004, 21:46:16; Search time 1.20792 Seconds (without alignments) 7673.534 Million cell updates/sec
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/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
               GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-252-991A-10598
US-09-620-312D-287
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US-09-433-826B-282
US-09-604-287A-282
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US-08-746-789A-1
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Result

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                 880, App
4285, Ap
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11508, A
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APPLICANT: Yuqui, Jiang
APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.470C3
CURRENT APPLICATION NUMBER: US/09/389, 681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FREESEQ for Windows Version 3.0
10256,
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; Sequence 282, Application US/09620405B
; Patent No. 6528054
; GENERAL INFORMATION:
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan L.
; APPLICANT: Harlocker, Susan L.
; APPLICANT: Harlocker, Susan L.
; APPLICANT: Hopler, William T.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
; FILE REFERENCE: 210121.470c8
; CURRENT FILING DATE: 2000-07-20
; NUMBER OF SEQ ID NOS: 495
; SOFTWARE: FRAESE for Windows Version 3.0
; SEQ ID NO 282
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100.0%; Pred. No. 0.92;
tive 0; Mismatches 0; Indels
                                US-09-252-991A-4285
US-09-702-705-1668
US-09-705-457-1668
US-09-129-033-1
US-09-129-033-1
US-09-512-991A-15054
US-09-512-91A-16
US-09-252-991A-856
US-09-252-91A-856
US-09-0461-775-8
US-09-041-166-8
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US-09-252-991A-815
US-08-461-775-10
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US-09-031-606-10
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RESULT 3
US-09-339-338-282

i Sequence 282, Application US/09339338A

; Patent No. 6573368

i GENERAL INFORMATION:

; APPLICANT: Widtu, Jiang

; APPLICANT: Mitcham, Jennifer L.

; TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND

; TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE

; TITLE PEPTERECE 2114.4702.

; CURRENT APPLICATION NUMBER: US/09/339,338A

; CURRENT FILING DATE: 1999-06-23

; NUMBER OF SEQ ID NOS: 315

; SOFTWARE: FRAESEQ for Windows Version 3.0
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APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILLE REFERENCE: 21011.470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 474
SOFTWARE: FastSEQ for Windows Version 3.0
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Patent No. 6579973
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Dillon, Davin C.
Mitcham, Jennifer L.
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; ORGANISM: Homo sapiens
US-09-433-826B-282
                        ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-620-405B-282
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LENGTH: 502
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1 CCGGGACATCCTCATCCACCC 21

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GENERAL INFORMATION:
APPLICANT: McClelland, Michael
APPLICANT: Welsh, John
TITLE OF INVENTION: Using Same
TITLE OF INVENTION: Using Same
FILE REFERENCE: P-PH 3457
CURRENT APPLICATION NUMBER: US/09/300,958A
CURRENT APPLICATION NUMBER: US/09/301
PRIOR PILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1998-04-27
PRIOR FILING DATE: 1999-08-27
PRIOR PLICATION NUMBER: 60/098,070
PRIOR FILING DATE: 1999-08-27
PRIOR PLICATION DATE: 1999-02-04
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                                                                                                                                                                                                            APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Jillon, Davin C.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Withouth Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
FILE REPRERENCE: 210121.470C7
CURRENT APPLICATION NUMBER: US/09/604,287A
CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 21; DB 4; Length 502; 100.0%; Pred. No. 0.92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                       ; Sequence 282, Application US/09604287A; Patent No. 6586572
290 CCGGGACATCCTCATCCACC 310
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; Sequence 27, Application US/09300958A
; Patent No. 6495319
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SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
....has 21; Conservative
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Best Local Similarity 100.(
Matches 21; Conservative
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US-09-604-287A-282
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; ORGANISM: Homo sapiens
US-09-300-958A-27
                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                              RESULT 5
US-09-604-287A-282
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LENGTH: 1907
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; LOCATION: (96)...(1211)

; OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)

; OTHER INFORMATION: protein.

US-09-570-593-4
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Patent No. 5789200

GENERAL INFORMATION:

APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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100.0%; Score 21; DB 4; Length 1907;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 21; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                          APPLICANT: Kaufmann, Joerg
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Marrow. Gree
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REPRESENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270 5219
TELEPHONE: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                               Sequence 4, Application US/09570593
Patent No. 6566063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                          GENERAL INFORMATION:
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LENGTH: 1907
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US-08-746-789A-1
US-09-570-593-4
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US-09-328-352-99/c
US-09-328-352-99/c
US-09-328-352-99/c
Sequence 99, Application US/09328352
Patent No. 6562968
GENERAL INFORMATION:
APPLICANT'S GATY L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT PILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
LENGTH: 1086
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Pred. No. 1.1e+02;
0; Mismatches 1; Indels 0;
                                                                                                                                                                Query Match 100.0%; Score 21; DB 1; Length 1920; Best Local Similarity 100.0%; Pred. No. 1; Aatches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lemke Ph.D. et al., Greg E.
TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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CLASCIPICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/237,401
FILING DATE: 02-MAY-1994
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US 07/884,486
FILING DATE: 15-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/456,647B
FILING DATE: 02-UN-1995
CLASSIPICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                               951 CCGGGACATCCTCATCCACCC 971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/08456647B
Patent No. 5811516
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                1 CCGGGACATCCTCATCCACCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Acinetobacter baumannii
US-09-328-352-99
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Best Local Similarity 94.4%;
Matches 17; Conservative
                    TYPE: Nucleic Acid
STRANDENNESS: Single
TOPOLOGY: Linear
JANTL-SENSE: No
US-08-746-789A-1
LENGTH: 1920
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US-08-456-647B-1/c
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ZIP: 92037
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us-08-978-217-13.rni

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TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 US-09-643-597-303
                                    ; NAME/KEY:
; LOCATION:
US-08-237-401A-1
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77.1%; Score 16.2; DB 1; Length 165;
Best Local Similarity 85.7%; Pred. No. 1.1e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-237-401A-1/C

Sequence 1, Application US/08237401A

Sequence 1, Application US/08237401A

Patent No. 5837441

APPLICANT: Lemke Ph.D. et al., Greg E.

TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES

NUMBER OF SEQUENCES: 54

CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson P.C.

STREET: 4225 Executive Square, Suite 1400

CITY: La Jolla

STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/237,401A
FILING DATE: OMAY-1994
CLASSIPICATION: 435
PRICIS APPLICATION NUMBER: US/08/4486
FILING DATE: IS-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Haile Ph.D., Lisa A.
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: UNIMBER: US/0251/007001
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell Ph.D., John R.
REGISTRAINON NUMBER: 31.678
REFERENCE/DOCKET NUMBER: 07251/007002
TELECOMMUNICATION INFORMATION:
TELEFAX: (6.9) 678-5070
TELEFAX: (6.9) 678-5070
TELEFAX: (6.9) 678-5070
INFORMATION FOR SEQ ID NO: 1.
SEQUENCE CHARACTERISTICS:
LENGTH: 165 base pairs
TYPE: nucleic acid
STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CCGGGACATCCTCATCCACCC 21
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TELEPAN: (619) 678-5070
TELEPAN: (619) 678-5099
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 165 Base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA
IMMEDIATE SOURCE:
CLONE: Tyro-1
                                                                                                                                                                                                                                                                                                                                                                                                                     ; NAME/KEY: CDS
; LOCATION: 1..165
US-08-456-647B-1
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CLONE: Tyro-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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Gaps
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Query Match
77.1%; Score 16.2; DB 2; Length 165;
Best Local Similarity 85.7%; Pred. No. 1.1e+02;
Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Li, Samuel X.
APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Robert A.
APPLICANT: McNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Hosken, Nancy A.
APPLICANT: Kalos, Michael D.
APPLICANT: Kalos, Michael D.
APPLICANT: Fanger, Gary R.
APPLICANT: Fanger, Cary R.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C6
CURRENT APPLICATION NUMBER: US/09/480,884A
CURRENT FILING DATE: 2001-08-27
NUMBER OF SEQ ID NOS: 330
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 210121.455C11
CURRENT APPLICATION NUMBER: US/09/643,597
CURRENT FILING DATE: 2000-08-21
NUMBER OF SEQ ID NOS: 369
SOFTWARE: FASTESEQ for Windows Version 3.0
SEQ ID NO 303
LENGTH: 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-480-884A-303
Sequence 303, Application US/09480884A
Setter No. 6482597
GENERAL INFORMATION:
APPLICANT: Wang, Tongtong
APPLICANT: Fall, Liqun
                                                                                                                                                                                                                                                                                                     Sequence 303, Application US/09643597
Patent No. 6426072
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                                                                                                               1 CCGGGACATCCTCATCCACCC 21
                                                                                                                                                                 76 ccedercarcerchageaces se
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; CTHER INFORMATION: n = A,T,C or G
US-09-643-597-303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bangur, Chaitanya S.
Hosken, Nancy
Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                       Wang, Tongtong
Fan, Liqun
Kalos, Michael D.
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NAME/KEY: misc_feature
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77.1%; Score 16.2; DB 4; Length 361;
Best Local Similarity 85.7%; Pred, No. 1.2e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                      Query Match 77.1%; Score 16.2; DB 4; Length 361; Best Local Similarity 85.7%; Pred. No. 1.2e+02; Matches 18; Conservative 0; Mismatches 3; Indels (
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mang, Tongtong
APPLICANT: Pan, Liqun
APPLICANT: Pan, Liqun
APPLICANT: Bangur, Chaiteanya S.
APPLICANT: Bangur, Chaiteanya S.
APPLICANT: Hosken, Nancy A.
APPLICANT: Hosken, Nancy A.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REPERENCE: 210121.455C8
CURRENT APPLICATION NUMBER: US/09/542,615A
CURRENT APPLICATION NUMBER: US/09/542,615A
CURRENT FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 350
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 303, Application US/09606421B Patent No. 6531315
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 303, Application US/09542615A Patent No. 6518256
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                                                                                                                           NAME/KEY: misc_feature

LCCATION: (1)...(361)

CTHER INFORMATION: n = A,T,C or G

US-09-480-884A-303
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APPLICANT: Wang, Tongtong
APPLICANT: Ranos, Michael D.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Fanger, Gary R.
APPLICANT: Fanger, Gary R.
APPLICANT: Li, Samuel X.
APPLICANT: Mang, Aijun
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SEQ ID NO 303
LENGTH: 361
TYPE: DNA
ORGANISM: Homo sapien
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ORGANISM: Homo sapien
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US-09-606-421B-303
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LENGTH: 361
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APPLICANT:
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Sequence:

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Sequence Sequence

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Sequence 937, Application US/09016434
| Patent No. 6500938
| GENERAL INFORMATION:
| APPLICANT: Jafice Au-Young
| APPLICANT: Jafice Au-Young
| APPLICANT: Jafice Au-Young
| TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
| TITLE OF INVENTION: PATHWAY GENE EXPRESSION
| NUMBER OF SEQUENCES: 149
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
| STREET: 3174 POPTER DRIVE
| CITY: PALO ALTO
| STATE: CALIFORNIA
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILING DATE: HEREWITH
CLASSTFICATION NUMBER:
PRICR APPLICATION NUMBER:
FILING DATE:
FIL
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US-09-620-312D-226
US-08-537-400-15
US-08-706-702-17
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US-09-238-471-17
US-09-702-705-95
US-09-736-457-95
US-08-484-661A-38
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US-09-257-584-6
US-08-484-661A-7
US-08-484-661A-15
US-08-484-661A-15
US-08-484-661A-12
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PCT-US96-09641-38
US-08-484-661A-36
US-08-656-664-36
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US-08-484-661A-10
US-08-656-664-10
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US-08-484-661A-32
                                                                       US-09-146-053-4
US-07-741-453A-57
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NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0002 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                    RESULT 1
US-09-016-434-927
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Command line parameters:
-MODEL=frame+ p2n.model -DEF=x1h
-Q=/Cgn2_1/USPTO_spool/US08978217/runat_10022004_133826_20481/app_query.fasta_1.1500
-Q=/Cgn2_1/USPTO_spool/US08978217/runat_10022004_133826_20481/app_query.fasta_1.1500
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=biosum62 -TRANS=human40.cdi
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-UNITS=bits -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXEN=200000000
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-NO_MMAP -LANGEQUERY -NGG_SCORES=0 -MAIT -DSPBLOCK=100 -LONGLOG
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-FGAPEXT=7 -YGAPDP=10 -YGAPEXT=0.5 -DELDEXT=7
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Sequence 25, Appl
                                                                                                                                       February 12, 2004, 19:34:21; Search time 4.03329 Seconds (without alignments) 1750.959 Million cell updates/sec
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Sequence 1, Al
Sequence 25, 1
Sequence 5, Al
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Sequence 4, 7
Sequence 1, 7
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Sequence 1,
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1: /cgn2_6/prodate_1/ina/5A_COMB.seq:*
2: /cgn2_6/prodate_1/ina/5B_COMB.seq:*
3: /cgn2_6/prodate_1/ina/6A_COMB.seq:*
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5: /cgn2_6/prodate_1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/prodate_1/ina/PcTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd
                                                                                            OM protein - nucleic search, using frame_plus_p2n model
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US-09-300-958A-27
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US-08-746-789A-1
US-09-638-649-6
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US-07-920-281C-1
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US-09-299-141-5
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Maximum Match 100%
Listing first 45 summaries
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 200000000
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84
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Match Length DB
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Jatabase :

Result

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US-09-100-958A-27
US-09-100-958A-27

US-09-100-958A-27

Patent No. 649319

Patent No. 649319

GENERAL INFORMATION:

APPLICANT: Wcclelland, Michael

APPLICANT: Wcclelland, Michael

APPLICANT: Trenkle, Thomas

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: P-PH 3457

CURRENT FILING DATE: 1999-04-27

PRIOR FILING DATE: 1999-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 85

SONGWARE: PatentIn Ver. 2 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1161 AAAAACTCAAGCGCTGCAAGGAGGAAGAGGTTCTCCCAGAGTCGGAAC 1208
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Sequence 4, Application US/09570593

Sequence 4, Application US/09570593

Sequence 4, Application US/09570593

GENERAL INFORMATION:

APPLICANT: Kaufmann, Joerg

APPLICANT: Nan, Farses

FILE REFERENCE: 2300-1556

CURRENT FILING DATE: 2000-05-12

PRIOR APPLICATION NUMBER: 60/134,112

PRIOR APPLICATION NUMBER: 60/134,112

PRIOR APPLICATION NUMBER: 60/134,112

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FastSEQ for Windows Version 4.0

FEMALE APPLICATION NOME APPLICATION APPLICATION NOME APPLICATION APPLICATION APPLICATION APPLICATION APPLICATION NOME APPLICATION AP
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Mismatches:
Indels:
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Best Local Similarity:
Query Match:
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LOCATION: (96)
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Pred. No.:
Score:
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LENGTH: 1907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: MCICELLand, Michael
APPLICANT: WCICELLand, Michael
APPLICANT: WCICELLand, Michael
APPLICANT: WCICELLAND
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: UGing Same
FILE REFERENCE: P-PH 3457
CURRENT APPLICATION NUMBER: US/09/300,958A
CURRENT FILING DATE: 1999-04-27
PRIOR APPLICATION NUMBER: GO/083,331
PRIOR APPLICATION NUMBER: GO/083,030
PRIOR FILING DATE: 1998-04-27
PRIOR APPLICATION NUMBER: GO/098,070
PRIOR FILING DATE: 1998-06-27
PRIOR FILING DATE: 1998-02-04
NUMBER OF SEQ ID NOS: 85
SOFTWARE: PARCET NOS: 85
SOFTWARE: PARCET NOS: 85
TOWN 14.5
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Matches:
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Matches:
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Mismatches:
Indels:
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Sequence 25, Application US/09300958A
Patent No. 6495319
GENERAL INFORMATION:
             TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 927:
SEQUENCE CHARACTERISTICS:
LENGTH: 237 base pairs
TYPE: nucleic acid
STRANTEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.00%
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100.00%
100.00%
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CLONE: 773734
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similarity:
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; LOCATION: (458)
US-09-300-958A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: unsure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similari
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                                                                                                                                                                                                                                                                                                                                         US-09-016-434-927
                                                                                                                                                                                                                                                                                                                                                                                                       Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match:
DB:
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us-08-978-217-12.rni

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APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1180 AAAACTCAAGCGGCTGGAAGGAGGAAGGAGGTTCTCCAGAGTCGGAAC 1227
                                                                                                                                                                                                   1161 AAAAACTCAAGCGCTGGAAGGAGGAGGATCTCCCAGAGTCGGAAC 1208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 LysAsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                                                                                                          1 LysAsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 19406-0939
COMPUTER READABLE PORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COMPUTER: MINDOWS FOR WORKGROUPS
SOFTMARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
Indels:
         Conservative:
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                         Mismatches:
Indels:
                                                                                                                US-08-978-217-12 (1-16) x US-09-570-593-4 (1-1907)
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKilne Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                      Sequence 1, Application US/08746789A Patent No. 5789200 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-638-649-6/c
Sequence 6, Application US/09638649
Partent No. 6563015
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: William T. Han
REGISTRATION NUMBER: 34,344
REFRENCE/DOCKET NUMBER: ATG
TELEPCOMMUNICATION INFORMATION:
TELEPRAX: 610 270 5219
TELEPRAX: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.00%
      100.00%
100.00%
100.00%
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STRANDENESS: Single
TOPOLOGY: Linear
JANTI-SENSE: NO
US-08-746-789A-1
Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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Query Match:
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                                                                                                                                                                                                                                               RESULT 5
US-08-746-789A-1
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COUNTRY:
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APPLICANT: Stern, David M.
APPLICANT: Schmidt, Ann Marie
APPLICANT: Schmidt, Ann Marie
APPLICANT: Yan, Shi Du
TITLE OF INVENTION: TRANSGENIC MICE OVER-EXPRESSING RECEPTOR FOR ADVANCED
TITLE OF INVENTION: GLYCATION ENDPRODUCT (RAGE) AND MUTANT APP IN BRAIN AND
TITLE OF INVENTION: USES THEREOF
FILE REPERENCE: 0575/62175
CURRENT APPLICATION NUMBER: US/09/638,649
CURRENT APPLING DATE: 2000-08-14
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 boo.vo=-os-t-s)

betent No. 6475780

patent No. 6475780

general INPORMATION:
    APPLICANT: Parrington, Mark
    APPLICANT: Parrington, Mark
    APPLICANT: Li, Xiaomao
    IIIE REPLICANT: Michel H.
    ITILE OF INVENTION: ALPHAVIRUS VECTORS FOR PARAMYXOVIRUS VACCINES
    FILE REFERENCE: 1038-1042 MIS
    CURRENT APPLICATION NUMBER: US09/554,337
    CURRENT FILING DATE: 2000-10-20
    PRIOR PILING DATE: 1997-11-14
    PRIOR RILING DATE: 1997-11-14
    PRIOR RILING DATE: 1998-11-13
    NUMBER OF SEQ ID NOS: 7
    SOFTWARE: PatentIN Ver. 2.1
    SEQ ID NO 4
    LENGTH: 8100
    TODE: DATE: DATE
    TODE: DATE
    SEQ ID NOS: 7
    LENGTH: 8100
    TODE: DATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 SerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative:
Mismatches:
Indels:
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Matches:
Conservative:
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Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 275 AATTCCAGTGGCTGGAAGGAGGAG
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; ORGANISM: respiratory syncytial virus
US-09-554-337-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 AsnSerSerGlyTrpLysGluGlu 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 1, Application US/07920281C; Patent No. 5739026
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100.00%
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54.76%
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Best Local Similarity:
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Murine
                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Pate
SEQ ID NO 6
LENGTH: 1348
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Pred. No.:
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Pred. No.:
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US-08-466-277-1
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/note= "Semliki Forest Virus complete nucleotide
sequence, presented as a cloned DNA sequence; see
Figure 5."
       APPLICANT: Liljestrom, Peter
APPLICANT: Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on TITLE OF INVENTION: Alphaviruses
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          675 GCCACAAACTGGGCCGACGAGGTGTTACAGGCCAGGAAC 716
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                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/920,281C

FILING DATE: 13-AUG-1992

CLASSIFICATION: 435

ATTONREY/AGENT INFORMATION:

NAME: WHIPHY JT., Gerald M.

REGISTRATION NUMBER: 28,977

REFRENCE/DOCKET NUMBER: 828-103P

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 7421..11179
OTHER INFORMATION: /product= "SFV polyprotein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: 87..7379
OTHER INFORMATION: /product= "SFV polyprotein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-978-217-12 (1-16) x US-07-920-281C-1 (1-11517)
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Matches:
                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIF: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Semliki Forest Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 703-2-1
TELEX: 248345
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 11517 base pairs
TVPE: nucleic acid
TVPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: RNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46.00
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7421..11179
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LOCATION: 1..11517
OTHER INFORMATION: /
OTHER INFORMATION: /
OTHER INFORMATION: /
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Best Local Similarity:
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alignment Scores:
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RESULT 9

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NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STRATE: Vigalia
CONDUTER: Vigalia
CONDUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BAP PC compatible
COMPUTER: BAP C compatible
COMPUTER: COMPUTER: PATOLIA Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: Us/08/466,277
FILING DATE: CONFORM:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 28,977
TELEFRAX: 703-241-1304
TELEFRAX: 703-241-2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: -
LOCATION: 1..11517
OCHER INFORMATION: /label= genome
/note= "Semliki Forest Virus complete nucleotide sequence, presented as a cloned DNA sequence; see Figure 5."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: CDS
LOCATION: 7421..11179
COTHER INFORMATION: /product= "SFV polyprotein"
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-08-466-277-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: CDS
LOCATION: 87..7379
OTHER INFORMATION: /product= "SFV polyprotein"
Sequence 1, Application US/08466277
Patent No. 6190666
GENERAL INFORMATION:
APPLICANT: Garoff, Henrik
APPLICANT: Garoff, Henrik
Illjestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Semliki Forest Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID No: 1:
SEQUENCE CHARACTERISTICS:
TYPE: nucleic acid
STRANDEDRESS: single
TOPOLOGY: linear
MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46.00
85.71$
50.00$
54.76$
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Best Local Similarity:
Query Match:
DB:
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Query Match:
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                                                                                                                                                                                                                                                                             Sequence 1, Application US/09554337

Sequence 1, Application US/09554337

Patent No. 647580

GENERAL INFORMATION;

APPLICANT: Barington, Mark

APPLICANT: Li, Xiaomao

ITILE OF INVENTION: ALPHAVIRUS VECTORS FOR PARAMYXOVIRUS VACCINES

FILE REFERENCE: 1038-1042

CURRENT APPLICATION NUMBER: US/09/554,337

CURRENT FILING DATE: 1997-11-14

PRIOR APPLICATION NUMBER: PCI/CA98/01064

PRIOR FILING DATE: 1997-11-14

PRIOR FILING DATE: 1998-11-13

NUMBER OF SEQ ID NOS: 7

SEQ ID NO: 1

SEQ ID NO: 1
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                                                                                                                                                    675 GCCACAAACTGGGCCGACGAGCAGGTGTTACAGGCCAGGAAC 716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 SerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                    3 SerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
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Requence 25, Application US/09464535
Retent No. 6542200
GENERAL INFORMATION:
APPLICANT: Famodu, Omolayo O.
APPLICANT: Caboon, Rebecca E.
APPLICANT: Caboon, Rebecca E.
APPLICANT: McGonigle, Brian
APPLICANT: McGonigle, Brian
APPLICANT: McGonigle, Brian
APPLICANT: Mafalski, J. Antoni
TITLE OF INVENTION: STEROL BIOSYNTHETIC ENZYMES
FILE REFERENCE: BB1306 US NA
CURRENT APPLICATION NUMBER: US/09/464,535
CURRENT PILING DATE: 1999-12-15
EARLIER APPLICATION NUMBER: 60/112,555
EARLIER FILING DATE: 1998-12-16
NUMBER OF SEQ ID NOS: 44
SOPTWARE: Microsoft Office 97
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Conservative:
Mismatches:
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Mismatches:
Indels:
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    US~08-978-217-12 (1-16) x US-08-466-277-1 (1-11517)
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , ORGANISM: respiratory syncytial virus US-09-554-337-1
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64.29%
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85.71%
50.00%
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CORGANISM: Glycine max
US-09-464-535-25
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Best Local Similarity:
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Best Local Similarit
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Sequence 12, Application US/08060925A
Patent No. 543924
GENERAL INFORMATION:
PATENT: Brantley, Mark
APPLICANT: Laubach, Victor
TITLE OF INVENTION: INTROPESSION OF ALPHA-1
TITLE OF INVENTION: INTROPIN IN EXPRESSION VECTORS THROUGH THE INCLUSION OF TITLE OF INVENTION: INTRO II
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 AsnSerSerGlyTrp------LysGluGluGluValLeuGlnSerArg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence:p43C-AT-INUS-09-299-141-5
                                                                                                                                                                                                                                         APPLICANT: FLOTTE, TERENCE R.
APPLICANT: SONG, SIHONG
APPLICANT: SONG, SIHONG
APPLICANT: MORGAN: MICHAEL
TITLE OF INVENTION: MATERIALS AND METHODS FOR GENE THERAPY
FILE REFERENCE: 4300,011800
CURRENT FILING DATE: 1999-04-23
EARLIER APPLICATION NUMBER: 60,083,025
EARLIER APPLICATION NUMBER: 60,083,025
EARLIER PILING DATE: 1998-04-24
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN Ver. 2.0
                                                                                       661 rccaccaccearccaarcaarcaaccararracaarccacc 620
                                           2 AsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArg 15
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620 NEWPORT CENTER DRIVE SIXTEENTH FLOOR
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Conservative:
Mismatches:
Indels:
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US-08-978-217-12 (1-16) x US-09-464-535-25 (1-993)
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                                                                                                                                                         US-09-299-141-5/c; Sequence 5. Application US/09299141

Patent No. 6461606; GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45.00
61.90%
47.62%
53.57%
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Best Local Similarity:
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US-08-060-925A-12/c
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LENGTH: 7492
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6394 AACAGCTCAGGCTGGTTGAGCAACCTTAACAAGAAGATGTAATTCACCAGAGCAAA 6335
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APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Zhang, Jie
APPLICANT: Mang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Xu, Chongjun
APPLICANT: Xu, Chongjun
APPLICANT: Xu, Chongjun
APPLICANT: Xu, Chongjun
APPLICANT: No. 6436703el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 790CIP25.
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
FRIOR APPLICATION NUMBER: 2000-03-31
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PL-Genes Version 2.0
LENGTH: 2763
                             SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,925A
FILING DATE: 06-MAY 1993
CLASSIFICATION: 435
ATTONENY/AGENT INFORMATION:
NAME: FUller, Michael L.
REGISTRATION NUMBER: 36,516
REGISTRATION FORSEY ID NO: 12: 12 ELEPAX: 619-235-0176
INFORMATION FOR SEQ ID NO: 12: 5EQUENCE CHARACTERISTICS: 12: 16-235-0176
INFORMATION FOR SEQ ID NO: 12: 5EQUENCE CHARACTERISTICS: 12: 16-235-0176
INFORMATION FOR SEQ ID NO: 12: 5EQUENCE CHARACTERISTICS: 12: 16-235-0176
SEQUENCE CHARACTERISTICS: 15: 16-235-0176
STRANDENSS: 31: 10: 16-235-0176
STRANDENS
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Mismatches:
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Matches:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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Sequence 8, Application US/09668680
Patent No. 6436703
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Liu, Chenghua
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-060-925A-12
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2 - AsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
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Matches:
Conservative:
Mismatches:
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                            44.00
66.67%
53.33%
52.38%
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44.00
66.67%
53.33%
52.38%
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Liu, Chenghua
, LOCATION: (180)..(1742)
US-09-668-680-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: (27)...(2375)
                                                                                                       Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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Best Local Similarity:
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Pred. No.:
                                                         Alignment Scores:
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us-08-978-217-7.rni

Page

Sequence 7550, Ap Sequence 7788, Ap Sequence 12113, A Sequence 12252, A Sequence 11100, A Sequence 10964, A

Sequence 3, Appli Sequence 78, Appl Sequence 1, Appli Sequence 1, Appli

Sequence

Sequence 7, Appli Sequence 1, Appli Sequence 37, Appl Sequence 7864, Ap Sequence 7864, Ap Sequence 7380, Ap Sequence 7245, Ap

1, Appli 1, Appli 9, Appli

Sequence Sequence Sequence Sequence 516, App Sequence 10, Appl

Sequence 20, Sequence 20, equence 20,

Sequence 13, Appli Sequence 1, Appli

App1

Sequence 9,

Sequence:

Run on:

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RESULT 1

US-09-100-956A-27

Sequence 27, Application US/09300958A

Patent No. 6495139

GENERAL INFORMATION:

APPLICANT: Mccllland, Michael

APPLICANT: Mccllland, Michael

APPLICANT: Trenkle, Thomas

TILLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TILLE REFERENCE: P-PH 3457

CURRENT FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-02-04

NUMBER OF SEQ ID NOS: 85

SOFTWARE: PATENTIN VICENCE: 1998-02-04
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US-09-338-933-107
US-08-215-681-107
US-08-316-6918-23
US-09-167-206-3
US-09-167-206-3
US-09-167-206-3
US-09-351-217-1
US-09-351-215-1
US-09-351-215-1
US-09-352-991A-7788
US-09-352-991A-7788
US-09-352-991A-7788
US-09-352-991A-7864
US-08-952-991A-7864
US-08-952-991A-7864
US-08-952-991A-7864
US-08-952-991A-7864
US-09-252-991A-7864
US-09-252-991A-7455
US-08-162-809-13
US-09-252-991A-7455
US-08-162-809-13
US-09-252-991A-7455
US-08-162-809-13
US-09-222-903-371A-20
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Mismatches:
Indels:
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Matches:
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: HOTH
US-09-300-958A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alignment Scores:
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LENGTH: 1907
                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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MODEL=frame+_p2n.model -DEV=xlh

MODEL=frame+_p2n.model -DEV=xlh

-0=/cgn2_1/USR970 spool/USR970217/runat_10022004_133826_20481/app_guery.fasta_1.1500
-DB=18sued_Patents_NA -OFWT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -STRAR=1 -END=-1 -TMATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-USER=LGCAL -OUTPWT=pto -NORMeaxt - HEAPSIZE=500 -MINLENS 0 -MAXLENS=200000000
-USER=US08978217 @CGN 1 1.115 @runat_10022004_133926_20481 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPELOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 27, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 13629, A
Sequence 13819, A
Sequence 139, App
Sequence 139, App
Sequence 92, Appl
                                                                                                                                                      February 12, 2004, 19:34:21; Search time 21.1748 Seconds (without alignments) 1750.959 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1 NCALEELRLVFGFLGDQLHA......BLLDDGQQASPYHFGSCGAG
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1: /cgn2_6/ptodatcal/lina/5A_COMB.seq:*
/cgn2_6/ptodatcal/lina/5B_COMB.seq:*
3: /cgn2_6/ptodatcal/lina/6A_COMB.seq:*
/cgn2_6/ptodatcal/lina/6B_COMB.seq:*
5: /cgn2_6/ptodatcal/lina/PCTUG COMB.seq:*
6: /cgn2_6/ptodatcal/lina/PCTUG COMB.seq:*
                        GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                                        OM protein - nucleic search, using frame_plus_p2n model
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US-09-570-59-4
US-09-476-789A-1
US-09-252-991A-13629
US-09-252-991A-13629
US-09-404-879A-139
US-09-318-913-139
US-09-215-681-139
US-09-215-681-139
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US-09-215-681-139
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US-09-215-681-139
US-09-215-681-22
US-09-216-68-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   569978 seqs, 220691566 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUMMARIES
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 200000000
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Match Length DB
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                                                                                                                                                                                                                                                   Title:
Perfect score:
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445
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Result

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645 rerececación 656
            81 CysGlyAlaGly 84
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STRANDEDNESS: Single
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; ANTI-SENSE: NO
US-08-746-789A-1
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Best Local Similarity:
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                                                                             RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41 GluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 GlnLeuArgAspleuThrSerSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeu 40
                                                                                                                                                                                                                          21 GlnLeuArgAspLeuThrSerSerSerSerAspGluLeuSerTrplleIleGluLeuLeu 40
                                                                                                                                    41 GluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer 60
                                                                                                                                                                                                       61 ProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySer 80
1 ASECYSALALEUGLUGLULEUARGLEUVALPHEGLYPROLEUGLYASPGINLEUHISALA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (BSX)
OTHER INFORMATION: protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KIN, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REPRESENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09570593
Patent No. 6566063
GENERAL INFORMATION:
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Best Local Similarity:
Query Match:
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LOCATION: (96)
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LENGTH: 1907
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41 GlurysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer 60
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585 CCCITTGCCCAGGAGCTGCTGGACGACGGTCAGCAAGCCAGCCCCTACCACCCCGGCAGC 644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla
                                                                                                                                                     US-08-746-789A-1

Sequence 1, Application US/08746789A

Petent No. 5789200

PERENT INFORMATION:

APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck

APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck

TITLE OF INVENTION:

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: SmithKline Beecham Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIF: 19406-0939
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COMPUTER: MICHOSOFT WORD
SUFFRATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746, 789A
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                               SEE: SmithKline Beecham Corporation: 709 Swedeland Road, P.O. Box 1539 King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: William T. Han
REGISTRATION UNDRER: A134,344
REFERENCE/DOCKET NUMBER: ATG 50024
TELEPHONE: 610 270 5219
TELEPHONE: 610 270 4026
INFORMATION SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILLING DATE:
ATTORNEY/AGENT INFORMATION:
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273 ACTCTCATCAGCAGCTTTTTCAGC-----TTCTTCCAGCTTTTGCAGGGC 229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 GluLeuArgleuValPhe-----GlyProLeuGlyAspGlnLeuHisAlaGlnLeu 22
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                                                                                                                                                                                                                                                                                                                                               32 GlubeuSerTrplleIleGlubeuLeuGlubysAspGlyMetAlaPheGlnGluAlabeu 51
                                                                                                                                                                                                                                                                                                                                                                                                                                52 AspproGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGln 71
                                                                                                                                                                                                                                                                 12 GlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThrSerSerSerSerAsp 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            333 GAGTICCAICTITCTICAICTITIAAGGCCCGGITITICAAIAACCTICATACCTCTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Mitchen, Jennifer L.
APPLICANT: Ming, Gordon E.
APPLICANT: Ming, Gordon E.
APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: 110400015 OF OVARIAN CANCER
FILE REFERENCE: 210121.462C2
CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT PILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              72 GlnAlaSerProTyrHisProGlySerCysGlyAlaGly 84
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Matches:
Conservative:
Mismatches:
Indels:
                                                                          Length:
Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 139, Application US/09404879A Patent No. 6468546 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (1)...(521)
OTHER INFORMATION: n = A,T,C or
, ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-13819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.61
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Best Local Similarity:
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                                                                                                                         Percent Similarity:
Best Local Similari
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REPERENCE: 1096.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR PAPLICATION NUMBER: US 60/074,788
PRIOR PILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
NUMBER OF SEQ ID NOS: 33142
LENGTH: 801
                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT:
MARC JO, RUDERIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ALERGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE REFERENCE: 107196.136
TITLE REPERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR PLILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF EQ. ID NOS: 33142
SEQ. ID NO 13629
LENGTH: 624
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Matches:
Conservative:
Mismatches:
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Patent No. 6551795
GENERAL INFORMATION:
                                                                                                                                                                     Sequence 13629, Application US/09252991A Patent No. 6551795
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Query Match:
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273 ACTCTCATCAGCAGCTTTTTCAGC-------TTCTTCCAGCTTTTGCAGGGC 229
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                                                    ----ProPheAspGlnGlySerProPheAlaGlnGluLeuLeu 67
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Patent No. 6488931

GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
FILE REFERENCE: 210121.462c1
CURRENT APPLICATION NUMBER: US/09/338,933
CURRENT APPLICATION NUMBER: US/09/338,933
CURRENT PILING DATE: 1999-06-23
NUMBER OF SEC ID NOS: 312
SOFTWARE: PastSEQ for Windows Version 3.0
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Matches:
Conservative:
Mismatches:
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LOCATION: (1)...(521)
OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
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LENGTH: 521
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273 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGCTTTTGCAGGGC 229
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APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: Algare, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REPERENCE: 210121.46502
CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
                                                                                              APPLICANT: Mitchan, Jennifer L.
APPLICANT: Frudakis, Tony N.
APPLICANT: Frudakis, Tony N.
APPLICANT: Frudakis, Tony N.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: OF OVARIAN CANCER
FILE REFERENCE: 210121.463
CURRENT APPLICATION NUMBER: US/09/215,681A
CURRENT PILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 310
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 139
LENGTH: 521
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Mismatches:
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Matches:
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                                       Sequence 139, Application US/09215681A Patent No. 6528253 GENERAL INFORMATION:
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US-09-404 879A-92/c
1G-09-404 879A-92/c
Sequence 92, Application US/09404879A
Parent No. 6468546
GENERAL INFORMATION:
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; OTHER INFORMATION: n = A,T,C or G
US-09-215-681-139
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ORGANISM: Homo sapien
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Best Local Similarity:
RESULT 8
US-09-215-681-139/c
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294 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGGTTTTTGCAGGGC 250
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354 GAGTICCAICTITICITCAICTITIAAGGCCCGGITTICAAIAACCTICAIACCTCTCTC 295
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                                   23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLys 42
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Patent No. 652823

GENERAL INFORMATION:

APPLICANT: Mitcham, Jennifer L.

APPLICANT: King, Gordon B.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: OF COMPOSITIONS AND METHODS FOR DIAGNOSIS

CURRENT APPLICATION NUMBER: US/09/215,681A

CURRENT FILIGATION NUMBER: 1999-12-17

NUMBER OF SEQ ID NOS: 310

SOFTWARE: FASTSEQ for Windows Version 3.0
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Matches:
Conservative:
Mismatches:
Indels:
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US-09-215-681-92
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US-09-215-681-92/c
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LENGTH: 551
TYPE: DNA
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394 ACTCTCATCAGCAGCTTTTTCAGC-----TTCTTCCAGCTTTTTGCAGGC 250
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APPLICANT: Ming, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
FILE REPERENCE: 210121.462C1
CURRENT APPLICATION NUMBER: US/09/338,933
CURRENT PILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 312
SOFTWARE: PastSEQ for Windows Version 3.0
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Indels:
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Matches:
   SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 92 LENGTH: 551
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Patent No. 6488931
GENERAL INFORMATION:
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US-09-338-933-92
                                                                              ORGANISM: Homo sapien
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LENGTH: 551
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; SEQ ID NO 107
; LENGTH: 555
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-338-933-107
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Sequence 107, Application US/09404879A

Patent No. 6468546

GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer L.
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121-462C2
CURRENT APPLICATION NUMBER: US/09/404,879A

CURRENT PILLING DATE: 1999-09-24

NUMBER OF SEQ ID NOS: 393

SOFTWARE: FREESEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
TITLE OF INVENTION: OVARIAN CANCER
CURRENT APPLICATION NUMBER: US/09/338,933
UUMBER OF SEQ ID NOS: 312
SOFTWARE: FASELSEQ for Windows Version 3.0
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Mismatches:
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Patent No. 6488931
GENERAL INFORMATION:
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US-09-404-879A-107
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Best Local Similarity:
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DB:
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358 GAGTICCATCITICITCATCITITAAGGCCCGGITITCAATAACCTTCATACCTCTC 299
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Patent No. 652853
GENERAL INPORMATION:
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Frudakis, Tony N.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
TITLE REFERENCE: 210121.463
CURRENT PILLING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 310
SOFTWARE: PARLSEQ for Windows Version 3.0
                555
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                                                Conservative:
Mismatches:
Indels:
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                  Length:
Matches:
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                                                 Percent Similarity:
Best Local Similarity:
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US-09-215-681-107/c
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Alignment Scores:
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LENGTH: 555
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388 CAAGGAGCCACCTCAGCCTGTTCCCGGGCCCGCCTTTCTCCCT-CAACTTCTC 330
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                                                                    493 ACTCICATCAGCAGCTTTTTCAGC------TTCTTCCAGCTTTTGCAGGC 449
                                                                                                                                                                                                                                                                        448 AGTGGCCAGGCGCCCCGAGGCACGGTCCAGCTCTTCAACCAGCTGGATCCTACGGTT 389
                                                                                                                                                  23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrplleIleGluLeuLeuGluLys 42
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Gaps:
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  358 GAGTICCAICTITICTICAICTITIAAGGCCCGGITITICAAIAACCTICAIACCTCTCTC 299
                                                                             -------TTCTTCCAGCTTTTGCAGGGC 254
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                                                                                                                                                                                                                                                                                                                       134 GCTGGAGGCGCTCAGCTCCTCTCTCTCTCTGCCTGCTGCAGAACCTGGATCT 75
                                        23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLys 42
                                                                                                                        43 AspGlyMetAla------54
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APPLICANT: Calabretta, Bruno
APPLICANT: Skorski, Tomasz
TITLE OF INVENTION: ANTESNSE
TITLE OF INVENTION: OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
TITLE OF INVENTION: OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
NUMBER OF SEQUENCES: 55
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seidel, Gonda, Lavorgna & Monaco, P.C.
STREET: TWO Penn Center, Suite 1800
CITY: Philadelphia
STATE: Pennsylvania
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Mismatches:
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MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
COMPUTER: IEM PS/2
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/306,691B
FILING DATE: September 15, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Monaco, Daniel A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 8321-8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-5849
TELERA: No. 5734039e
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 2301 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23, Application US/08306691B Patent No. 5734039
                                                                           298 ACTCTCATCAGCAGCTTTTTCAGC-
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OPERATING SYSTEM: MS-DOS
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TYPE: nucleic acid
STRANDEDNESS: double
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TGCGCTTCA 66
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COUNTRY: U.S.A.
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Best Local Similarity:
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US-08-306-691B-23
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TYPE: DNA
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Sequence 3437, Ap
Sequence 12009, A
Sequence 12009, A
Sequence 8097, Ap
Sequence 1, Appli
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Sequence 4, Appli
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9092, Ap
9180, Ap
8895, Ap
8997, Ap
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Sequence 4, Appli
                                                                 February 12, 2004, 21:46:16; Search time 14.4951 Seconds (without alignments) 7673.534 Million cell updates/sec
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                                                                                                                                      1 AATIGIGCCCTIGAGGAGCT...........CCGGCAGCIGIGGCGCAGGA 252
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Sequence 8
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/cgm2_6/ptodata/1/ina/5B_COMB.seq:*
/cgm2_6/ptodata/1/ina/6A_COMB.seq:*
/cgm2_6/ptodata/1/ina/6A_COMB.seq:*
/cgm2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgm2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-242-677-1
US-08-252-991A-12006
US-09-252-991A-12006
US-09-252-991A-8097
US-09-252-991A-8097
US-09-1252-991A-8097
US-09-185-370-1
US-09-111-444-4
US-09-111-444-4
US-09-552-991A-9180
US-09-252-991A-9180
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US-09-252-991A-7739
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                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
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US-09-570-593-4
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                                                                                                                                                                                                 569978 segs, 220691566 residues
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Maximum Match 100%
Listing first 45 summaries
                                            OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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252
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Match Length
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Perfect score:
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ALIGNMENTS

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Sequence 27, Application US/09300958A

Patent No. 649519

GENERAL INFORMATION:

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: Trenkle, Thomas

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Using Same

FILE REFERENCE: P-PH 3457

CURRENT APPLICATION NUMBER: US/09/300,958A

CURRENT APPLICATION NUMBER: 60/083,331

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

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Best Local Similarity
US-09-300-958A-27
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OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
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Patent No. 5789200
Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELP3
NUMBER OF SEQUENCES:
ADDRESSEE: SMITCHAINE Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 252; DB 4; Length 1907; 100.0%; Pred. No. 5e-59; tive 0; Mismatches 0; Indels 0
                                                                                         APPLICANT: Kaufman, Joerg
APPLICANT: Kaufman, Joerg
APPLICANT: Kin, Hong
APPLICANT: Alexewe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
TITLE OF INVENTION: CANCER
TITLE OF INVENTION: CANCER
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: 2300-1556
CURRENT APPLICATION NUMBER: 60/134,112
PRIOR PELLING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
                         Sequence 4, Application US/09570593
Patent No. 6566063
GENERAL INFORMATION:
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US-09-570-593-4
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APPLICANT: Laigudi, Raghunath V.
APPLICANT: Laigudi, Raghunath V.
APPLICANT: Laigudi, Raghunath V.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR FILE REFERENCE: PL-0017 US.
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
SEQ THANKE: PERL PROGRAM
SEQ ID NO 3437
LENGTH: 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 484 CAGCTGCGAGACTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG
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; OTHER INFORMATION: Incyte ID No. 6476212 700611585H1
US-09-313-294A-3437
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY AGENT INFORMATION:
NAME: William T. Han
REGISTRATION NUMBER: 34,344
REGISTRATION NUMBER: ATT 50024
TELECOMMUNICATION INFORMATION:
TELEFONDEY 610 270 5219
TELEFONDES: 610 270 5219
TELEFONDES: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 3437, Application US/09313294A; Patent No. 6476212; GENERAL INFORMATION:
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Matches 252; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
NTI-SENSE: No
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; ANTI-SENSE:
US-08-746-789A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
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Sequence 12009, Application US/09252991A
Sequence 12009, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ARRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR PLING DATE: 1999-02-18
PRIOR PLING DATE: 1999-02-18
PRIOR PLING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 12066
LENGTH: 741
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             149 gecesarceccesarcascersescrescariaciaciaciaciaciateaacecaaceccaacecceas 208
                                                                             108 CATTGAGCTGCTGGAAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTT 167
                                                                                                                                                                                                                                         193 CCACCTCGCGCGCGCCTCCGGGAGCGCCCTGCGCCCCCCGCTGCCGCGCGCCTCCTCGT 134
                                                                                                                                                           133 cerceadececricadaadaaaacecadecercecacecercecececarecerecerr 74
                                                                                                                                                                                                                                                                                                                  16 GAGCIGCGICIGGICITIGGGCCICIGGGGGACCAACICCAIGCCCAGCIGCGAGACCIC
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14.4%; Score 36.2; DB 4; Length 741;
Best Local Similarity 47.6%; Pred. No. 0.51;
Matches 107; Conservative 0; Mismatches 118; Indels (
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US-09-252-991A-12066
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US-09-252-991A-12066
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and Uses Thereof in regulating Gene Expression and in the
Treatment of AIDS
                                                                                                                                                                        42 GGGGGACCAACTCCATGCCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAG 101
                                                                                                                                                                                                                                                      60. Geredadeceacrecededadercereceaceerrerecearerreceareceerrece 119
                                                                                                                                                                                                                                                                                                                                        102 TIGGATCATIGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGG 161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   162 GCCCTTTGACCAGGCAGCCCCTTTGCCCAGGAGCTGCTGGACGACGGTCAGCAAGCCAG 221
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                                                                                               0; Gaps
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    Query Match 15.5%; Score 39; DB 4; Length 276; Best Local Similarity 49.3%; Pred. No. 0.07; Matches 102; Conservative 0; Mismatches 105; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/242,677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   222 CCCCTACCACCCGGCAGCTGTGGCGC 248
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ATTONEX/AGENT INFORMATION:
NAME: Mayfield, Denise L.
REGISTRATION NUMBER: 33,732
REPREBUCE/DOCKET NUMBER: UTSD:401
TELECOMMUNICATION:
TELEPHONE: 713-787-1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433 CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-242-677-1/c
; Sequence 1, Application US/08242677
; Patent No. 5677143
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APPLICANT: Gaynor, Richard B
APPLICANT: W., Foon W.
TITLE OF INVENTION: Cellular Nu
TITLE OF INVENTION: and Uses Th
TITLE OF INVENTION: Treatment o
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              713-789-2679
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LOCATION: 1..4863
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48 CCAACTCCATGCCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGAT 107

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Sequence 8205, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION:
NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
FILE REFERENCE: 107196.136
CURRENT FILING DATE: 1999-02-18
PRIOR PILING DATE: 1999-02-18
PRIOR PILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 8205
LENGTH: 2145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1576 CCAGCCCTGGGCGCCCCGCGTCCACCGGCGATCTCGGCGGCGGCGAGGCACAGGCGGCGGAA 1635
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APPLICANT: Kyriakis, John M.
APPLICANT: Pombo, Celia M.
APPLICANT: Bonyontre, Joseph
TITLE OF INVENTION: SOK-1 AND METHODS OF USE
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/852,743
FILING DATE: 7-MAY-1997
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/016,774
FILING DATE: 7-MAX-1996
ATTORNEY/AGENT INFORMATION:
NAME: FTRSET, Janis K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 1, Application US/08852743
; Patent No. 5830699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 104; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sest Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-252-991A-8205
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US-08-852-743-1
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Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
SENERAL INFORMATION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
CURRENT PELICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     676 GAGCGCTTCCTCGCCGCCGCGCAGCGCCCTGGAACTGGACCTGCGCCTGCTGTACCTC 735
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14.4%; Score 36.2; DB 4; Length 1650;
Best Local Similarity 47.9%; Pred. No. 0.62;
Matches 104; Conservative 0; Mismatches 113; Indels 0
                                                                                                                                                                                                                                                                      14.4%; Score 36.2; DB 4; Length 918; 47.6%; Pred. No. 0.53;
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                                                                                                                                                                                                                                                                                                                                                   0; Mismatches 118; Indels
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                                                                                                                                                     ORGANISM: Pseudomonas aeruginosa
        NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 107; Conservative
                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                         US-09-252-991A-12009
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US-09-252-991A-8097
                                        SEQ ID NO 12009
LENGTH: 918
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                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                          Query Match
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RESULT 9 US-09-252-991A-8205

0; Gaps

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355 racarcaccecraciriescre 377
 TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                     INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 1975 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                          Coding Sequence 127...1404
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                                                                                                                                                                                                                                                                                                                                       98; Conservative
                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: CDNA
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Best Local Similarity
Matches 98; Conserv
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                                                                                                                                                                                                                        ; NAME/KEY:
; LOCATION:
US-09-185-370-1
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                                                                                                                                                                                                                                                                                                                                         13.9%; Score 35; DB 2; Length 1975; 48.3%; Pred. No. 1.4; tive 0; Mismatches 105; Indels
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APPLICANT: Kyriakis, John M.
APPLICANT: Pombo, Celia M.
APPLICANT: Bonventre, Joseph
TITLE OF INVENTION: SOK-1 AND METHODS OF USE
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COPERATING SYSTEM: Windows95
SOFTWARE: PastSRQ for Windows Version 2.0
CURBENT APPLICATION DATA:
APPLICATION NUMBER: US/09/185,370
FILING DATE:
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00786/327001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTER STICS:
LENGTH: 1975 base pairs
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/852,743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09185370 Patent No. 6093560 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                  Coding Sequence 127...1404
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                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 48.33
Matches 98; Conservative
                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                            MOLECULE TYPE: CDNA
                                                                                                                                                                                                              linear
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                                                                                                                                                                                                                                                                NAME/KEY:
LOCATION:
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46 GACCAACTCCATGCCCAGCTGCGAGACCTCCACTTCCAGCTCTTCTGATGAGCTCAGTTGG 105
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13.9%; Score 35; DB 3; Length 1975; 48.3%; Pred. No. 1.4;
                                                                                                                                                                              0; Mismatches 105; Indels
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Patent No. 5863780
GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION NUMBER: US/08/712,709
FILING DATE: Filed Herewith
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0118 US
TELECOMMUNICATION INPORMATION:
TELEPHONE: 415-855-0555
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
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STRANDEDNESS: single
TOPOLOGY: linear
  CLONE: Consensus
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Best Local Similarity
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MOLECULE TYPE:
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ZIP: 94304
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                                                                                                                                                                                              0; Gaps
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APPLICANT: Au-Young, Janice
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
INTHER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94304
COMPUTER READBELE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COPERATING SYSTEM: DOS
SOFTWARE: FESTESQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/111,444
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APPLICATION NUMBER: 08/712,709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/09111444 Patent No. 6045792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
STRANDEDNESS: single
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Best Local Similarity
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                    linear
                                                                                            CLONE: Consensus
US-08-712-709-4
              TOPOLOGY: lines
MOLECULE TYPE: CL
IMMEDIATE SOURCE:
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IMMEDIATE SOURCE:
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Best Local Similarity 4813;

Best Local Similarity 48
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RESULT 15

US-09-222-991A-8781/C

SEQUENCE SET 1911, Application US/09252991A

PACTOR NO. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND ANINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPBUTICS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPBUTICS

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 8781

LENGTH: 621
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13.8%; Score 34.8; DB 4; Length 621;
Best Local Similarity 54.8%; Pred. No. 1.2;
Matches 69; Conservative 0; Mismatches 57; Indels
0; Mismatches 105; Indels
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ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-8781
     Matches 98; Conservative
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Sequence 1, A sequence 2, A sequence 2, A sequence 1, A sequence 1, A sequence 1, A sequence 3, A sequence 1, A se

Sequence 1 Sequence 1 Sequence 1

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RESULT 1
US-09-300-958A-27

Sequence 27, Application US/09300958A

Patent No. 6495319

GENERAL INFORMATION:
APPLICANT: Wclelland, Michael
APPLICANT: Welsh, John
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Using Same
TITLE OF INVENTION: Using Same
FILE RFERENCE: P-PH 3457

CURRENT APPLICATION NUMBER: US/09/300,958A

CURRENT APPLICATION NUMBER: 60/098,070

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-09-27

PRIOR FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 85

SOFTWARE: Patentin Ver: 2.0

SECTION OF SEQ ID NOS: 85
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Matches:
Conservative:
Mismatches:
                US-09-439-313-44

US-09-352-616A-44

US-09-032-616A-44

US-09-039-913-8

US-09-009-913-8

US-09-009-913-8

US-09-009-913-10

US-09-009-913-10

US-09-05-913-10

US-09-105-113-2

US-09-570-593-12

US-09-570-593-12

US-09-570-593-12

US-09-570-593-12

US-09-570-593-12

US-09-354-579-13

US-09-320-759-10

US-09-920-759-11

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ORGANISM: Homo sapiens
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Sequence 4, Appli
Sequence 1, Appli
                                                                                                                                              February 12, 2004, 19:34:21; Search time 93.522 Seconds (without alignments) 1750.959 Million cell updates/sec
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                                                                                                                                                                                                                                                                                 1 MAATCEISNIFSNYFSAMYS.....YKFGKNSSGWKEEEVLQSRN 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/ptogrus_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/packfiles1.seg:*
                   GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                        - nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-300-958A-27
US-09-570-593-4
US-08-746-799A-1
US-09-399-681-282
US-09-393-882-82
US-09-333-86-282
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US-09-009-913-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 569978 seqs, 220691566 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                         BLOSUM62
Kgapop 10.0 , Kgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                         US-08-978-217-2
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Matches:
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   1176 TGGAAGGAAGAAGATTCTCCAGAGTCGGAAC 1208
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                                                                                                     RESULT 2
US-09-570-593-4
; Sequence 4, Application US/09570593
; Patent No. 6566063
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORM
US-09-570-593-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1056 AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 1115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1116 ATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGC 1175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             936 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCCATGAAGTGGGGAAATCGGCATGAA 995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | IleLeuGluArgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGly 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      221 ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         756 CCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGGGGATCCC 815
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           301 GlyValPheLygPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys 320
                                                      96 ATGGCTGCAACCTGTGAGATTAGCAACATTTTTAGCAACTACTTCAGTGCGATGTACAGC 155
                                                                                                                                                                                                              215
                                                                                                                                                                                                                                                                                                                               216 GTACTGACCCTGAGCAACCCCCAGATGTCATTGGAGGGTACAGAGAAGGCCAGCTGGTTG 275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GluLysAsnLysTyrAspAlaSerAlaIleAspPheSerArgCysAspMetAspGlyAla 100
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MetAlaAlaThrCysGluIleSerAsnIlePheSerAsnTyrPheSerAlaMetTyrSer 20
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                                                                                                                                    SergluAspSerThrLeuAlaSerValProProAlaAlaThrPheGlyAlaAspAspLeu
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                                                                                                                                                                                                                                                                                        41 ValLeuThrLeuSerAsnProGlnMetSerLeuGluGlyThrGluLysAlaSerTrpLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp
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81 GluLysAsnLysTyrAspAlaSerAlaIleAspPheSerArgCysAspMetAspGlyAla 100
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LOCATION: (96)...(1211)
OCHER INPORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
APPLICANT: Kaufmann, Joerg
APPLICANT: Kin, Hong
APPLICANT: Xin, Hong
APPLICANT: Azrowe, Greg
APPLICANT: Azrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REFERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FRASEG for Windows Version 4.0
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Patent No. 5789200
GENERAL INFORMATION
GENERAL INFORMATION: A No. 5789200el Human BTS Family Member, ELF3
TITLE OF INVENTION: A No. 5789200el Human BTS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Smithkline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Frussia
STATE: PA
COUNTRY: USA
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ZIP: 19406-0939

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
CORPATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MCROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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Matches:
                            REGISTRATION NUMBER: 34,344

REGISTRATION NUMBER: 34,344

REFERENCE/DOCKET NUMBER: ATG 50024

TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELEFRAX: 610 270 4026

INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1920
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
TOPOLOGY: Linear
NATI-SENSE: NO
US-08-746-789A-1
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Query Match:
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US-09-620-405B-282
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US-09-620-405B-282
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  241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 260
                                     835 AAGCACGGGAAGCGGAAACGAGGCCGGCCCCGAAAGCTGAGCAAAGAGTACTGGGACTGT 894
                                                                             261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280
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APPLICANT: Viqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470c3
CURRENT APPLICATION NUMBER: US/09/389,681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTI: 502
                                                                                                            895 CTCGAGGGCAAGAAGAAGAAGAACACGCGCCCAGAGGCACCCACCTCCTGTGGGAGTTCATCCGG
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8.09-389-681-282
Sequence -282, Application US/09389681A
Patent No. 6518237
GENERAL INFORMATION
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                                                                                                             243 AAGAAGAGACAAGCACCCCAGAGGCACCCACTGTGGAGTTCATCCGGGACATCCTC 302
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183 AAGCGGAAACGAGCCCGCCCCCGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGGC 242
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                                                                      LysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArgAspIleLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillom, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Mariocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND ITLLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.47008
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 4958
SOFTWARE: FREUSEQ for Windows Version 3.0
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303 ATCCACCGGGAGCTCAACGAGGGCCTCATGAAGTGGGGAAATCGGCATGAAGGGCTCTTC 362
                                                                                                                                                    324 MetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrTysArgGlulleLeuGlu 343
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APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.470C4
CURRANT PAPLICANTON: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.470C4
CURRANT FILLING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 474
SSCOTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LINGTH: 502
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                                                                    LysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsn
                                                                                                          363 AAGTICCIGCGCICCGAGGCIGIGGCCCAACIAIGGGGCCAAAAGAAAAAAACAACAGCAAC
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Matches:
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; Patent No. 6579973
; GENERAL INFORMATION:
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Best Local Similarity:
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                                            243 AAGAAGAGCACGCGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCTC 302
                                                                                         284 IleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 303
                                                                                                                             303 ATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAAAATCGGCATGAAGGCGTCTTC 362
                                                                                                                                                                            304 LysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysBysAsnSerAsn 323
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                                                                                                                                                                                                                                                                                                           423 ATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGAA 482
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  264 LysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeu 283
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Batent No. 6573368
GENERAL INFORMATION:
APPLICANT: Vadiu, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: My Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 21012.470C2
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FastSEQ for Windows Version 3.0
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LENGTH: 502
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21-JAN-1998
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ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: RASISEQ for Win
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47.42%
36.08%
28.23%
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APPLICATION NUMBER: US,
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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STRANDEDNESS: double
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Best Local Similarity:
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US-09-009-913-2
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Sequence 282, Application US/09604287A

Sequence 282, Application US/09604287A

GENERAL INPORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470c7
CURRENT FILING DATE: 200-06-22

NUMBER OF SEQ ID NOS: 489

SOFTHARE: FREISEQ for Windows Version 3.0
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LENGTH: 502
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Matches:
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                                                                                                                                                 Sequence 2, Application US/09009913

Patent No. 6087485
GENERAL INFORMATION:
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES:
CRRESPONDENCE ADDRESS:
DDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                      SYSTEM: DOS FASTEMS Version 2.0
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315 TGGGAGTGGCTCCAGCACCTCCTGACCAGCTGGATGCCAATGTATCCTTTC 374
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------TrpLeuGlyGluGlnProGlnPheTrpSerLysThrGlnVal
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3-09-009-913-3 Sequence 3, Application US/09009913 Patent No. 6087485

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416 CTCCTGGACACCAACCAACCAGGTGCCAATTGTATCCCTTTCCAAGAGTTCGAACATCAAC 475
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Mismatches:
Indels:
                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
OPERATING SYSTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
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Matches:
GENERAL INFORMATION:
APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 21-JAN-1998
CLASSIFICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36,677
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
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SEQUENCE CHARACTERISTICS:
LENGTH: 5510 base pairs
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555.50
51.24%
39.13%
28.06%
                                                                                                                                                                                                                                                                                  IBM Compatible
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ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,
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MEDIUM TYPE: Diskette
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STRANDEDNESS: double
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                      734 CGGGCTCAGATCTCCATGACAACCACCAGTCACCTTCCTGTTGCAGAGTCACCTGATATG 793
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          284 IleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 303
176 AlaSerProTyrHisProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAsp 195
                                                                                         196 ValSerThrAlaGlyThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySer 215
                                                                                                                                   644 ---TCCATCATGAACACCTGGAAAGACGAGAACTATTTATATGACACCAACTATGGTAGC 700
                                                                                                                                                                                  216 AspvalAspleuAspProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCys 235
                                                                                                                                                                                                                            701 ACAĞTAĞATİTĞ-----TTGĞACAGCAAAACTİTĞ-------------------TĞC 733
                                                                                                                                                                                                                                                                              ------GlyAspProLysHisGly 243
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
FILLING DATE: 21-JAN-1998
CLASSIFICATION:
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Patent No. 6087485
GAPPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: BOZICAVIC & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                            635 ACTGAGCCT------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         196 ValSerThrAlaGlyThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySer 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             891 CGGGCTCAGATCTCCATGACAACCACCAGTCACCTTCCTGTTGCAGAGTCACCTGATATG 950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                244 LysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCysLeuGluGly 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 136 LeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetAlaPheGlnGluAlaLeuAsp 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             156 ProGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGln 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        792 ACTGAGCCT------ 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   216 AspValAspLeuAspProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCys 235
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        264 LysLysSerlysHisAlaProArgGlyThrHisLeuTrpGluPhelleArgAspIleLeu 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      513 CTCCTGGACACCAACCAGCTGGATGCCAATTGTATCCTTTCCAAGAGTTCGACATCAAC 632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  693 CAGCTCCTCTACAGCAACTTGCAGCATCTGAAGTGGAACGGCCAGTGCAGTAGTGAC--- 749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99 GlyalaThrieuCysAsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGly 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         59 TrpLeuGlyGluGlnProGlnPheTrpSerLysThrGlnValLeuAspTrpIleSerTyr 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 79 GlnValGluLysAsnLysTyrAspAlaSerAlaIleAspPheSerArgCysAspMetAsp 98
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                                                                                                                                                                                                                                                                                                           5667
126
39
78
79
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Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-978-217-2 (1-371) x US-09-009-913-4 (1-5667)
                                                                                                                                                                                                                                                                                                               Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                     Indels:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEQ
TELECOMMUNICATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              750 -----
                                                                                                                            INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 5667 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                               555.50
51.24*
39.13*
28.06*
                                                                                           TELEFAX: 650-327-3231
TELEX:
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                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                    Percent Similarity:
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GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
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                                                                                91 AspPheSerArgCysAspMetAspGlyAlaThrLeuCysAsnCysAlaLeuGluGluLeu 110
                                                                                                         790 CCTTTCCANGAGTTCGACATCAACGGGGGCACCTTTGCAGGATTGAGTTTGCAGGAGGTC 731
                                                                                                                                                                                                                                                 130 -----SerSerSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGly 147
                                                                                                                                                                                                                                                                                                                                       148 MetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGln 167
                                                                                                                                                                                                                                                                                                                                                                               ------ ACACAC 632
                                                                                                                                                                                                                                                                                                                                                                                                                       168 GluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySerCysGlyAlaGly 187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ------TCCATCATGAACACCTGGAAAGACNAGAACTAT 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             208 HisSerSerAspSerGlyGlySerAspValAspLeuAspProThrAspGlyLysLeuPhe, 227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----GlyAspProLysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLys 255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 256 GluTyrTrpAspCysLeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeu 275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             262 GGTAAAAAGAAGAAGAACAGCAGCATGACCTATGAAAAGCTCAGCCGAGCTATGAGAATAT 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        336 TyrTyrLysArgGluIleLeuGluArgValAspGlyArgArgLeuValTyrLysPheGly 355
                                                                                                                                                                 111 ArgleuValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThr--- 129
                                                                                                                                                                                                         730 ACCCGGGCGGCAGGGACGGCGGGGGCANCTCCTCTACAGCAACTTGCAGCATCTGAAGTGG 671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       276 TrpGluPhelleArgAspIleLeulleHisProGluLeuAsnGluGlyLeuMetLysTrp 295
                        71 GlnValLeuAspTrplleSerTyrGlnValGluLysAsnLysTyrAspAlaSerAlalle 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   565 TIATATGACACCAACTATGGTAGCACAGTAGATTTG-----TIGGACAGCAAAACTTTC
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                                                                                                                                                                                                                                                                                          670 AACGGCCAGTGCÁGTÁGTGÁC-----
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US-09-030-607-44/c
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US-09-020-956-44/c

US-09-020-956-44/c

Sequence 44, Application US/09020956

Patent No. 6261562

GENERAL INFORMATION:
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION:
CORRESPONDENCES: 178

CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP

STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                            1221 CGTGTGGATGGACGAAGACTGGTATATAAATTTGGGAAGAAGAAGAAGGATGGAGGATGGAGAAA 1280
                        .041 TTGAACCCAGACAAGAACCCAGGATTAATAAAATGGGAAGACCGATCTGAGGGGGGGTCTTC 1100
                                                                                                                      1101 AGGITCITGAAATCAGAGGGAGTGGCTCAGCTATGGGGTAAAAAGAAGAACAACAGCAGC 1160
                                                                                                                                                                                              1161 ATGACCTATGAAAAGCTCAGCCGAGCTATGAGATATTACTACAAAAAGAAAATACTGGAG 1220
                                                                              304 LysPheleuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsn 323
                                                                                                                                                                                                                                                    344 ArgValAspGlyArgArgArgLeuValTyrLysPheGlyLySAsnSerSerGlyTrpLysGlu 363
284 IleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 303
                                                                                                                                                               324 MetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGluIleLeuGlu 343
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
CLASSIFICATION:
ATTORNEY/AGRAT INFORMATION:
NAME: Maki, David J.
REFERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative:
Mismatches:
Indels:
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Matches:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEPAX: (206) 682-6031
INFERMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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519.50
50.97%
39.03%
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TYPE: nucleic acid
STRANDEDNESS: single
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MOLECULE TYPE: CDNA
ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                    364 GluGlu 365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      130 -----SerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGly 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       148 MetalaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGln 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     168 GluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySerCysGlyAlaGly 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         188 AlaProSerProGlySerSerAspValSerThrAlaGlyThrGlyAlaSerArgSerSer 207
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            CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
PILING DATE: US/09/030,607
FILING DATE: 25-FEB-1998
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 210121.427C3
TELECOMMUNICATION INFORMATION:
TELEPAK: (206) 622-4900
TELEPAK: (206) 632-6031
INFORMATION FOR SEQ ID NO: 444:
SEQUENCE CHARACTERISTICS:
                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
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519.50
50.97%
39.03%
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TYPE: nucleic acid
STRANDEDNESS: single
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NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                              CLASSIFICATION:
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                                                                                                 USA
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                                                                                               COUNTRY:
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208 HisSerSerAspSerGlyGlySerAspValAspLeuAspProThrAspGlyLysLeuPhe 227
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JAPPICANT: M. Jiangchun

JAPPICANT: Dillon, Davin C.

APPICANT: Mitcham, Jennifer L.

APPICANT: Mitcham, Jennifer L.

APPICANT: Harlocker: Susan Louise

APPICANT: Farlocker: Susan Louise

APPICANT: Ralos, Michael

APPICANT: Ralos, Michael

APPICANT: Reter, Mark

APPICANT: Reter, Mark

APPICANT: Reter, Mark

APPICANT: Bolk, John

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND

TITLE OF INVENTION: UNFORMER OF 120121.427C9

CURRENT APPLICATION NUMBER: 105/09/439,313

CURRENT FILING DATE: 1999-11-12

JUMBER OF SEQ ID NOS: 575

SOOTWARR: FastSEQ for Windows Version 3.0

LENGTH: 852

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Matches:
Conservative:
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LOCATION: (1)...(852)
OTHER INFORMATION: n = A,T,C or G
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Mismatches:
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US-09-352-616A-44/c
; Sequence 44, Application US/09352616A
Best Local Similarity: 39.03%
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                    GENERAL INFORMATION:
APPLICANT: Dillor, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jiangchun
APPLICANT: Mitcham, ComPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210.11.42708
CURRENT APPLICATION NUMBER: US/09/352,616A
NUMBER OF FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
SOFTWARE: FREUSEQ FOR Windows Version 3.0
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Matches:
Conservative:
Mismatches:
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OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
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Patent No. 6395278
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US-09-352-616A-44
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SEQ ID NO 44
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            316 GlyGlnLygLygLygLygAsnSerAsnMetThrTyrGluLygLeuSerArgAlaMetArgTyr 335
256 GluTyrTrpAspCysLeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeu 275
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-389-661-282
US-09-620-405B-282
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US-09-009-913-2
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US-09-016-434-927
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US-09-009-913-8
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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Sequence 307, App sequence 3, Appli Sequence 10, Appli Sequence 11, Appli Sequence 11, Appli Sequence 3, Appli Sequence 11, Appli Sequence 11, Appli Sequence 11, Appli Sequence 12, Appli Sequence 25, Appli Sequence 25, Appli Sequence 25, Appli Sequence 25, Appli Sequence 25, Appli Sequence 11, Appli Sequence 25, Appli Sequence 25, Appli Sequence 26, Appli Sequence 26, Appli Sequence 26, Appli Sequence 27, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli Sequence 28, Appli	Targets and Methods of	Length 1907; Lindels 0; Gaps 0; CTTCAGTGCGATGACAGC 60
US-09-016-434-307 US-09-920-759-3 US-09-920-759-10 US-09-36-779-1 US-09-435-335-1 US-09-435-335-1 US-09-435-344-579-1 US-09-016-434-840 US-08-343-443B-3 US-08-343-443B-3 US-09-016-048-3 US-09-021-715-1 US-09-021-715-1 US-09-021-715-1 US-09-920-759-11 US-09-920-759-11 US-09-620-312D-869	6 1 0 1 0 4	1.6e-273; es 0; TTAGCAACTA TTAGCAACTA TTAGCAACTA TGCTGCCAC
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2 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	RESULT 1 US-09-300-958A-27 Sequence 27, Appl Patent No. 649531 GENERAL INFORMATI APPLICANT: Wecl APPLICANT: Tren TITLE OF INVENTI; FILE REFERENCE: CURRENT PELING DATE PRIOR PELING DATE PERIOR PELING DATE PERIOR PELING DATE PERIOR PELING PERIOR PE	Query Match Best Local S Matches 1116 Qy 1 Db 96 Qy 61 Db 156

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 301 ACCCTCTGCAATTGTGCCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGGACCAA 360
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PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
                              Sequence 4, Application US/09570593
Patent No. 6566063
GENERAL INFORMATION:
US-09-570-593-4
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61 TCGGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCGATGACTTG 120
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                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: CDS
LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
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NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 1907
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Matches 1116; Conservative
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                                                                                                                                                                                                                                                                                                    AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 1020
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                                841 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 900
                                                                                                                                                                  GGCGTCTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCCAACTATGGGGCCAAAAGAAAAG 960
                                                                                             936 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1. Application US/08746789A
Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200e1 Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match

99.4%; Score 1109.6; DB 1; Length 1920;
Best Local Similarity 99.6%; Pred. No. 6.7e-272;
Matches 1112; Conservative 0; Mismatches 4; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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COMPUTER: IBM 486
COPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/746,789A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NOBER: US/08/746,789A
No. 5789200ember 15, 1996
N: 514
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REFERENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELEFAN: 610 270 4026
FORMATION PARTY 610 270 4026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: NO. 57892
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: William T. Han REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTI
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TYPE: Nucleic Acid
STRANDEDNESS: Singl
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HANTI-SENSE: NO
US-08-746-789A-1
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STATE: PA
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 3
US-08-746-789A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    122 TGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAGGGGGATCCCAAGCACGG 181
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                   Sequence 282, Application US/09389681A

Barent No. 6548237

GREERAL INFORMATION:
APPLICANT: Virgui, Jiang
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Ku, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 21011.47027
CURRENT APPLICATION NUMBER: US/09/389,681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
SOFTWARER: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   849 CATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44.7%; Score 499.4; DB 4; Length 502; 99.8%; Pred. No. 1.8e-117; vative 0; Mismatches 1; Indels 0
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Best Local Similarity 99.84
Matches 500; Conservative
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CORGANISM: Homo sapiens
US-09-389-681-282
US-09-389-681-282
                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 282
LENGTH: 502
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Sequence 282, Application US/09620405B Patent No. 6528054

RESULT 5 US-09-620-405B-282

GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangohun
APPLICANT: Hepler, William L.
APPLICANT: Hepler, William T.

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CATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGA 1028
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Sequence 282, Application US/09339338A

Fatent No. 6573368

GENERAL INFORMATION:
APPLICANT: Vugiu, Jiang
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: MITCHAM, Jangehm
TITLE OF INVENTION: INDRANOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
CURRENT APPLICATION NUMBER: US/09/339,338A

CURRENT PILING ADTE: 1999-06-23
NUMBER OF SEQ ID NOS: 315

SOSTWARE FESTSEQ for Windows Version 3.0
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAFY AND TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER FILE REFERENCE: 210121.47008
CURRENT APPLICATION WUMBER: US/09/620,405B
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 495
SOFTWARE: FASISEQ for Windows Version 3.0
SEQ ID NO 292
                                                                                                                                                                                                                                                                                                                                          44.7%; Score 499.4; DB 4; Length 502; 99.8%; Pred. No. 1.8e-117; cive 0; Mismatches 1; Indels 0
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Matches 500; Conservative
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US-09-339-338-282
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LENGTH: 502
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                                                       609 TTCTCGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTGGATCCCACTGA
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APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Jillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Wicham, Jennifer L.
APPLICANT: Wicham T.
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470C7
CURRENT APPLICATION NUMBER: US/09/604,287A
CURRENT FILING DAIT: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SEQ ID NOS: 882
LENGTH: 502
LENGTH: 502
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CRGANISM: Homo sapiens
US-09-604-287A-282
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
CURRENCE: 210121.470C4
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 474
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                                                                                                                                   Indels 0; Gaps
                                                                           Query Match 44.7%; Score 499.4; DB 4; Length 502; Best Local Similarity 99.8%; Pred. No. 1.8e-117; Matches 500; Conservative 0; Mismatches 1; Indels 0;
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Best Local Similarity 99.8%; Pred. No. 1.8e-117;
Matches 500; Conservative 0; Mismatches 1; Indels 0
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; Patent No. 6579973
; GENERAL INFORMATION:
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CRGANISM: Homo Bapiens
US-09-433-826B-282
ORGANISM: Homo sapiens
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     ; US-09-339-338-282
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LENGTH: 502
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GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS
NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
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                       818 CCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCGGGGCTCAACGAGGCCTCA 877
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APPLICATION NUMBER: US/09/030,607 FILING DATE: 25-FEB-1998 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
15.6%; Score 173.8; DB 3;
Best Local Similarity 69.7%; Pred. No. 6.5e-35;
Matches 235; Conservative 0; Mismatches 102;
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NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REPERENCE/DOCKET NUMBER: 210121.427C3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-030-607-44/c
; Sequence 44, Application US/09030607
; Patent No. 6262245
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INFORMATION FOR SEQ ID NO: 44:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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Patent No. 6261562
GENERAL INFORMATION:
APPLICANT: Dilin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                            729 GAAGCGGAAACGAGGCCGGCCCCGAAAGCTGAGCAAAGATACTGGGACTGTCTCGAAGG 788
                                                                                                                                                             122 TGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGGGGGGATCCCAAGCACGG 181
                                                                                      182 GAAGCGGAAACGAGGCCGGCCCCGAAAGCTGAGCAAAGAGTACTGGGGACTGTCTCGAGGG 241
                                                                                                                                 789 CAAGAAGAGCAAGCACGCGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCT
                                                                                                                                                                                                                       849 CATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTT
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15.6%; Score 173.8; DB 3; Length 852;
Best Local Similarity 69.7%; Pred. No. 6.5e-35;
Matches 235; Conservative 0; Mismatches 102; Indels 0
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COUNTRY: USA

COMPUTER: READALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORPUTER: PREFER PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: SEED and BERRY LLP 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1029 ACGGGTGGATGGCCGGCGACT 1049
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INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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nucleic acid
EDNESS: single
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938 AACTATGGGGCCAAAAGAAAAGAACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCA 997
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; Patent No. 6395278
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US-09-232-149A-44/c
; Sequence 44, Application US/09232149A
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NAME/KEY: misc_feature
LOCATION: (1)...(852)
OTHER INFORMATION: n = A,T,C or G
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APPLICANT: Dillon,
                                                                                                                                                                                                                                                                                                                               RESULT 12
US-09-352-616A-44/c
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450 TGAAAAAGGAGCAAGACCCCCTGCCAAGTGCCACACAAAAAGCACACCCGAGAGGGA 391
                                                                                                                                     390 crcacriaridadanircarccacaacarccicrigaaccaagacaagaaccaddarraa 331
                                                                                                                                                                                                                                     330 TAAAATGGGAAGACCGATCTGAGGGGGTCTTCAGGTTCTTGAAATCAGAGGCAGTGGCTC 271
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APPLICANT: XU, Jiangchun
APPLICANT: XU, Jiangchun
APPLICANT: Mitchan, Davin C.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Rados, Michael
APPLICANT: Redo, Michael
APPLICANT: Redo, Michael
APPLICANT: Reter, Mark
APPLICANT: Solk, John
APPLICANT: Boy, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
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Patent No. 6329505
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; LCCATION: (1)...(852)
; CTHER INFORMATION: n = A,T,C or G
US-09-439-313-44
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270 AGCTATGGGGTAAAAAGAAGAACAACAGCAGCATGACCTATGAAAAAGCTCAGCCGAGCTA 211
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APPLICANT: Billon, Davin C.
APPLICANT: Barlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REPERRACE: 1210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SRQ ID NOS: 472.
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FILING DATE: 21-JAN-1998
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                     GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.42706
CURRENT APPLICATION NUMBER: US/09/232,149A
CURRENT FILING DATE: 1999-01-15
SOFTWARE: FASELSEQ for Windows Version 3.0
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COMPUTER READBLE FORM:
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
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Patent No. 6087485
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
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STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature

LOCATION: (1)...(852)

// OTHER INFORMATION: n = A,T,C or G

US-09-232-149A-44
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ORGANISM: Homo sapien
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Patent No. 6465611
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US-09-009-913-2
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Best Local Similarity 69.7%; Pred. No. 1.1e-34;
Matches 235; Conservative 0; Mismatches 102; Indels 0;
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATURG SYSTEM: DOS
SOFTWARE: FastSEC for Windows Version 2.0
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TILLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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APPLICATION NUMBER: US/09/009,913
                                                                            FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEQ-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 3, Application US/09009913; Patent No. 6087485
                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLLOGY: linear
MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                  TELEFAX: 650-327-3231
                     PRIOR APPLICATION DATA: APPLICATION NUMBER:
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CLASSIFICATION:
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US-09-009-913-3
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Search completed: February 13, 2004, 01:54:40 Job time : 65.1925 secs

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APPLICANT: Karen GLATT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
                           Sequence 101, App
Sequence 563, App
Sequence 192, App
Sequence 1105, Ap
Sequence 1105, Ap
Sequence 217, App
Sequence 64, App
Sequence 853, App
Sequence 853, App
Sequence 853, App
Sequence 944, App
Sequence 244, App
Sequence 282, App
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Sequence 44, Appl
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Sequence 1953, Ap
Sequence 12, Appl
Sequence 3261, App
Sequence 203, App
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Sequence 44,
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15. US-10-097-340-74

16. US-09-964-824h-101

17. US-09-964-824h-101

17. US-09-964-824h-101

17. US-09-964-824h-102

17. US-09-922-117-1105

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Publication No. US20030087250A1
GENERAL INFORMATION:
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APPLICANT: Manjula GANNAVARAPU
APPLICANT: Sebastian HORRSCH
APPLICANT: Steve G. KOVATS
APPLICANT: Rachel E. MSYERS
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
APPLICANT: AMISEN
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Gordon B. MILLS
Robert C. BAST, Jr.
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Xumei ZHAO
Karen GLATT
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-MODBLE-frame+ p2n.model -DEV=xlh
-MODBLE-frame+ p2n.model -DEV=xlh
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-DE-VDISTE-I /USFO spool/US08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
-DB=Published Applications NA -QFMT=fastap -SUPFTX=rnpb -MINMATCH=0.1
-LOOPCIL -LOOPCIL - LOOPEXT=0 - UNITS=bits -START=1 - NNDTRIX=blosum62
-LOOPCIL -LOOPCIL - OUNTS=bits -START=1 - NNDTRIX=blosum62
-THR MAXINS-0 -ALIGN=15 -MODB=LOOPL -OUTPMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIENS-200000000 - USR=-US08978217 @CGN 1 1 690 @runat 1002204 133827_20514
-NCPUS - ICPUS - NO MMAP -LARGEQUERY -NNG GCRES=0 - WAIT -DSPBLOCK=100
-LONGLOG -DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAFOP=6 -FGAFEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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1. /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2. /cgn2_6/ptodata/1/pubpna/US06_NEW PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US06_DUBCOMB.seq:*
4. /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
5. /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
6. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
7. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
8. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
9. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
10. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
11. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
12. /cgn2_6/ptodata/1/pubpna/US09_NEW PUB.seq:*
13. /cgn2_6/ptodata/1/pubpna/US09_NEW PUB.seq:*
14. /cgn2_6/ptodata/1/pubpna/US09_NEW PUB.seq:*
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19. /cgn2_6/ptodata/1/pubpna/US10_NEW PUB.seq:*
                                  GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                           - nucleic search, using frame_plus_p2n model
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Maximum Match 100%
Listing first 45 summaries
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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and is derived by analysis of
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Maximum DB seq length: 200000000
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GENERAL INFORMATION:
APPLICANT: Horrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
FILE REFERENCE: 68290-73
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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NUMBER OF SEQ ID NOS: 593
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                                              200 ThralaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219
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                                                                                        696 ACTGGTGCTTCTCGGAGCTCCCACACTCCAGACTCCGGTGGAAGTGACGTGGACT
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Mismatches:
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SEQ ID NO 101
LENGTH: 1915
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US-09-964-824A-101
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US-09-964-824A-101
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Mismatches:
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Matches:
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR PLING DATE: 2001-09-26
PRIOR PELICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR FILING DATE: 2001-09-16
PRIOR PELING DATE: 2001-09-26
PRIOR PELING DATE: 2001-09-26
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86.79%
86.21%
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ORGANISM: Homo sapiens
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LENGTH: 1907
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Db 1200 TGGAAGAGGAAGAGTTCTCCAGAGTCGGAAC 1232	RESULT 3 US-09-964-824A-563 ; Sequence 563, Application US/09964824A ; Patent No. US20020102531A1	; GENERAL INFORMATION: ; APPLICANT: Horrigan Stephen ; TITLE OF INVENTION: Cancer Gene Determination and T ; TITLE OF INVENTION: Sets	; FILE REPERBNCE: 689290-73 ; CURRENT APPLICATION NUMBER: US/09/964,824A ; CURRENT FILING DATE: 2001-09-27 ; PRIOR APPLICATION NUMBER: US/60/236,033	PRIOR FILING DATE: 2000-09-28 PRIOR APPLICATION NUMBER: US/60/236,032 PRIOR FILING DATE: 2000-09-28 PRIOR APPLICATION NUMBER: US/60/236,028) FRIOR FILING DATE: 2000-09-28) NUMBER OF SEQ ID NOS: 583) SOFTWARE: Patentin version 3.0) SEQ ID NO 563	; LENGTH: 1915 ; TYPE: DNA ; ORGANISM: Homo sapiens US-09-964-824A-563	1,19e-189 1707.00	Percent Similarity: 92.72\$ Conservative: Best Local Similarity: 86.79\$ Mismatches: Query Match: 86.21\$ Indels: DB: 10	US-08-978-217-16_COPY_2_371 (1-370) x US-09-964-824A-5 1 AlaAlaThrCygGluileSerAsnValPheSerAsnTyrP	123 GCTGCAACTGTGAGATTAGCAACATTTTAG	Db 183 GAGGACTCCAGCCTCTGTTCCCCCTGCTGCCACCT	Qy 40 LeuThrLeuAsnAsnGlnGlnMetThrLeuGluGlyProG	Qy 60 GluargProGlnPheTrpSerLysThrGlnValleuGluT	000	Db 363 AAGAACAAGTACGACGCCATTGACTTCTCACGAT Qy 100 LeuCysSerCysAlaLeuGluGluLeuArgLeuValPheG	Db 423 CTCTGCAATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTT	Qy 120 HishadlnLeuArghspLeuThrSerAsnSerSerAspG	Oy 140 LeuLeuGluLysAspGlyMetSerPheGlnGluSerLeuG	Db 543 CTGCTGGAGAAGGATGGCCTTCCAGGAGGCCCTA- Ov 160 GlnGlvSevBroDbealaGlnGlnLenLenAsnAsnGlvA	600 CAGGGCAGCCCTTTGCCCAGGAGCTGCTGGACG
123 GCTGCAACCTGTGAGATTAGCAACATTTTTAGCAACTACCTTCAGTGCGATGTACAGCTGG 182	21 GluAspFroThrLeuAlaProAlaProProThrThrPheGlyThrGluAspLeuVal 39	40 LeuThileuAsnAsnGinGinMetThrLeuGluGlyProGluLysAlaSerTrpThrSer 59	60 GluargProGlnPheTrpSerLysThrGlnValLeuGluTrpIleSerTyrGlnValGlu 79	80 LysabnLysTyraspalaSerSerIleaspPheSerargCysasnMetAspGlyalaThr 99	100 LeuCysSerCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeu 119 ::	120 HisAlaGlnLeuArgAspLeuThrSerAsnSerSerAspGluLeuSerTrpIleIleGlu 139	140 LeuLeuGluLysAspGlyMetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAsp 159	160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyr 179	180 CysSerThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199 :::	200 ThrAlaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219 	LeuThrGluSerLysValPheProArgAspAspPheThrAspTyrLysLysGlyGluPro 23	780 CCCACTGATGGCAAGCTCTTCCCCAGGGATGGTTTTCGTGACTGCAAGAAGGGGGATCCC 839 240 LysHisGlylysArglysArgGlyArgProArglysLeuSerlysGluTyrTrpAspCys 259	GCCCCGAAAGCTGAGCAAAGAGTACTGGGACTG PPCATGGJVThvHistenTrnglubhaTlear	CTCGAGGGCAAGAAGGAGGCACCCCAGAGGCACCCACTGTGGGGGGTTCATCGG	280 AspleteullehisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 299 	GlyValPheLysPheLeuArgSerGl	1020 GGCGTCTTCAAGTTCCTGCGCTCCGAGGTGTGGGCCCAACTATGGGGCCAAAAGAAAAA 1079 320 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 339	AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAAC	340 IleLeuGluArgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGly 359 1140	TrpLysGluGluValGlyGluSerArgAsn 370
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                                                                                                          243 CTGACCCTGAGCAACCCCCAGATGTCATTGGAGGTACAGAGAAGGCCAGCTGGTTGGGG 302
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                                                                                                                                                                                                              183 GAGGACTICCACCTIGGCCTICTTTCCCCTTGCTGCCTTTTGGGGCCGATGACTTGGTA 242
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                                                                                                                                                                                                                                                                                                                                                60 GluArgProGlnPheTrpSerLysThrGlnValLeuGluTrpIleSerTyrGlnValGlu 79
                                                                                                                                                                                                                                                            40 LeuThrLeuAsnAsnGlnGlnMetThrLeuGluGlyProGluLysAlaSerTrpThrSer 59
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                                                                                     1 AlaAlaThrCysGluIleSerAsnValPheSerAsnTyrPheAsnAlaMetTyrSerSer
                                                                                                                                                                      21 GluAspProThrLeuAlaProAlaProPro---ThrThrPheGlyThrGluAspLeuVal
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                                                                                                            720 ACTGGTGCTTCTCGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGAACCTGGAT 779
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180 CysSerThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199
                                         660 CCCGGCAGCTGTGGCGCAGAGCCCCCTCCCCTGGCAGCTCTGACGTCTCCACCGCAGGG 719
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US-09-880-107-3420
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APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
TILLE OF INVENTION: Gene Expression Profiles in Liver Cancer: FILE REFERENCE: 44921-5028-WG.
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE PARCELL VOC. 2.1
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ORGANISM: Homo sapiens
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US-09-880-107-3420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FASCEEQ for Windows Version 4.0
SEQ ID NO 1105
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: APPLICANT: APPLICANT.
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
Clapper, Jonathan D.
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Sequence 192 Application US/099677AI
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturing Energy and Partical Conversion Number: US/09/967,768A
CURRENT APPLICATION NUMBER: US/60/236,109
PRICR APPLICATION NUMBER: US/60/236,034
PRICR APPLICATION NUMBER: US/60/236,111
PRICR FILING DATE: 2000-09-28
PRICR APPLICATION NUMBER: US/60/236,111
PRICR FILING DATE: 2000-09-28
PRICR APPLICATION NUMBER: US/60/236,111
1140 ATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1199
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SOFTWARE: PatentIn version 3.0
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ORGANISM: Homo sapiens
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SEQ ID NO 192
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                                                            320 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlu
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND LIABRED TO CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILIG DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: PSELSEQ for Windows Version 4.0
SEQ ID NO 1105
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Publication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: V., Jiangchun
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sinth, John A.
APPLICANT: Sinth, Carole L.
APPLICANT: Mang, Yuqiu
APPLICANT: Mang, Aijun
APPLICANT: Mang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Saith, Carole L.
APPLICANT: Gapper, Jonathan D.
APPLICANT: Gapper, Jonathan S.
APPLICANT: Panger, Gary R.
APPLICANT: Panger, Gary R.
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              US-09-922-217-1105
                                                    Alignment Scores:
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962 GACATCCTCATCCACCCGGAGCTCAACGAGGGCTCATGAAGTGGGAGAAATCGGGATGAA 1021
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305 GAACAGCCCCAGTTCTGGTCGAAGACGCAGGTTCTGGACTGGATCAGCTACCAAGTGGAG 364
                                          LysAsnLysTyrAspAlaSerSerIleAspPheSerArgCysAsnMetAspGlyAlaThr 99
                                                                                100 LeuCysSerCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeu
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PAL33PIR
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
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Publication No. US20040005579A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                US-08-978-217-16_COPY_2_371 (1-370) x US-10-264-049-756 (1-1956)
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Mismatches:
                                                                                                                                                                                                            Length:
Matches:
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PRIOR FILING DATE: 2001-06-07
PRIOR PELLON NUMBER: US 60/209,467
PRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SEGTHARE: Patentin Ver. 3.1
SEQ ID NO 756
LENGTH: 1956
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1707.00
92,72$
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                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
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384 AAGAACTACTACGATGGCGCCATTGACTTCTCACGATGTGACATGGGCGCCACC 443
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| Publication No. US20030235915A1
| GRNERAL INFORMATION
| APPLICANT: SPECHT, THOMAS
| APPLICANT: HINZMANN, BERND
| APPLICANT: HINZMANN, BERND
| APPLICANT: PILARSKY, CHRISTIAN
| APPLICANT: PILARSKY, CHRISTIAN
| APPLICANT: ROSENTHAL, ANDRE
| APPLICANT: ROSENTHAL, ANDRE
| APPLICANT: ROSENTHAL, ANDRE
| TITLE OF INVENTION: TUMORS
| TITLE OF INVENTION: TUMORS
| FILE REFERENCE: SCH-1763
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      260 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg
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TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE.REFERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT PILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR PILING DATE: 2000-03-08
PRIOR PILING DATE: 1999-03-12
NUMBER: OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF EQUIPMENT OF
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Matches:
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Patent No. US20020052308A1
GENERAL INFORMATION:
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86.79%
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; ORGANISM: Homo sapiens
US-09-925-301-207
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Best Local Similarity:
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                                                                                                                                                                                                                                                                     Conservative:
Mismatches:
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Matches:
CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
FRIOR APPLICATION NUMBER: 09/646,673
PRIOR FILING DATE: 2000-09-20
PRIOR PILING DATE: 1990-03-19
NUMBER OF SEQ ID NOS: 202
SOFTWARE: Patentin Ver: 2.1
SOFTWARE: 2269
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1458.50
93.71%
88.36%
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; ORGANISM: Homo sapiens
US-10-131-410-64
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Best Local Similarity:
Query Match:
DB:
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Pred. No.:
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrits, Heather
APPLICANT: Becits, Heather
APPLICANT: Becits, Heather
APPLICANT: Becits, Heather
APPLICANT: Becits, John A.
APPLICANT: Magher, Madeleine Joy
APPLICANT: Magher, John A.
APPLICANT: Mang, Yuqiu
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Mang, Aijun
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; Patent No. US20020076414A1
; GENERAL INFORMATION:
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Best Local Similarity:
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US-09-922-217-853/c
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131 SerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetSerPheGlnGlu 150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27 AAGCTGAGCAAAGAGTACTGGGACTGT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lodes, Michael J.
Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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US-10-025-380-853/c
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                                         151 SerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAsp 170
                                                                                 324 GCCCTA---GACCCAGGGCCCTTTGACCAGGCCAGCCCCTTTGCCCAAGGAGCTGCTGGAC
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384 TCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAG 325
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Sequence 853, Application US/09833263
Sequence 853, Application US/09833263
Sequence 853, Application US/09833263
GENERAL INFORMATION:
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Madeleine J.
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE FILE SETERENCE: 210121.471C12.
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEC ID NOS: 1093
SOFTWARE: FRALSEQ for Windows Version 3.0
                                                                                                                                                267 GACGGTCAGCAAGCCACCCCTACCACCCCGGCAGCTGTGGCGCAGGAGCCCCCTCCCCC
                                                                                                                      171 AspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerPro
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US-09-833-263-853
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LENGTH: 626
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171 AspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerPro 190
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                               131 SerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetSerPheGlnGlu
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                                                                                      71 LeuGluTrpIleSerTyrGlnValGluLysAsnLysTyrAspAlaSerSerIleAspPhe
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CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. US20020076414A1
GENERAL INFORMATION:
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APPLICANT: Lodes, Michael J.
APPLICANT: Beerist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
Clapper, Jonathan D.
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US-09-922-217-944
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APPLICANT: Wang, Aljun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
CURRENT FILING DATE: 2001-04-10
CURRENT FILING DATE: 2001-04-10
CURRENT FILING DATE: 2001-04-10
TYPER OF SEQ ID NOS: 1093
SOFTWARE: FeatSEQ for Windows Version 3.0
SEQ ID NO 944
LENGTH: 563
TYPE: DNA
CORGANISM: Homo sapien
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Search completed: February 13, 2004, 01:46:24 Job time : 599.817 secs

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Sequence 74, Appl Sequence 101, App Sequence 563, App Sequence 192, App Sequence 1105, App Sequence 756, App Sequence 207, App Sequence 207, App Sequence 64, App Sequence 1601, App Sequence 1601, App Sequence 1953, App
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Sequence 853, App
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-922-217-1105

US-09-922-217-1105

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-060-036-237	US-09-998-598-2216	10-060-036-3	3-2	9-604-287A-2	-339-338-	3-551-621-2	-124-805-2	0-007-805-2	0-076-622-2	1-998-598-3	.066-543-323	9-867-7	-922-217-9	9-833-263-94	.0-025-380-94	US-10-085-117-235	-10-305-720-92	9-893-238-	US-10-085-117-253	0-029-386-264	US-09-864-761-14745	-10-029-386-127	0-085-117-8	-10-004-113-4	-09-972-546	175-523~	-10-085-117-22	-10-085-117-2	US-10-229-834A-6	
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Sequence 74, Application US/10097340

Publication No. US20030087250A1

GENERAL INFORMATION:
APPLICANT: Only DERSCH
APPLICANT: Stave GANATYAR
APPLICANT: Stave G. KOMATS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
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APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WESTERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Rachel E. WEYERS
APPLICANT: Karen GLATT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION NUCLES: 120/10/097.340
CURRENT FILING DATE: 2001-03-14
FRIOR PELING DATE: 2001-03-14
FRIOR PELING DATE: 2001-03-14
FRIOR APPLICATION NUMBER: 60/276,025
FRIOR PELING DATE: 2001-03-14
FRIOR APPLICATION NUMBER: 60/216,026
FRIOR PELING DATE: 2001-03-14
FRIOR APPLICATION NUMBER: 60/216,026
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Patent No. US20020102531A1
GENERAL INFORMATION:
APPLICANT: Horizgan, Stephen
TITLE OF INVENTION: Sets
FILE OF INVENTION: Sets
FILE REPERBACE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1394 CICCOCACCCTCCTCTTGGAATTACAAGCCTGGGGGTTTGAAGCTGACTTTATAGCTGCA 1453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7457 AGACTGCAGGGACGGAGGGGACAGGGTTGTGTCCTCGGTACTTC---CTGGACTGCCTTC 7513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1570 GGAGCACCGTGATGGAGGAGGAGGAGGGGCTCCAGCACTTCTTCTGGACTGGCGTT 1629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7573 IGCTATA--AATATTCCAGGIGTATATAGAGAGCTATTTTTCTAAAGCATTTCCCCTCC 7630
                                                                                                                                                                                                                                                                                                6999 AGGTATTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
                                                                                                                                                                                                                                                                                                                                                                                                                        1155 TTTGGCAAAAACTCAAGCGGCTGGAAGGAGGAAGAGGGTTCTCCAGAGTCGGAACTGAGGG 1214
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1454 AGTGTATCTCCTTTTATCTGGTGCCTCCTCAAACCCAGTCTCAGACACTTAAATGCAGAC 1513
                                                                                                                                                                                                                                                                                                                                          1095 AGGTACTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAG
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                                                                                                                                                                                                              Query Match 2.8%; Score 216; DB 15; Length 1907;
Best Local Similarity 66.7%; Pred. No. 3.3e-54;
Matches 456; Conservative 0; Mismatches 195; Indels 33; Gaps
PRIOR APPLICATION NUMBER: 60/323,580 PRIOR FILING DAFF: 2001-09-19 NUMBER OF SEQ ID NOS: 363 SOFTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CIGCICIICICCACIGAGIGCIGG 7654
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                                                                                                                              TYPE: DNA
CORGANISM: Homo sapiens
US-10-097-340-74
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                                                                                      SEQ ID NO 74
LENGIH: 1907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.8%; Score 215.6; DB 10; Length 1915; Best Local Similarity 65.6%; Pred. No. 4.4e-54; Matches 449; Conservative 0; Mismatches 204; Indels 31; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7631 CTGCTCTTCTCCACTGAGTGCTGG 7654
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CURRENT FILING DATE: 2001-09-27
PRIOR PEDLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
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; Parent No. US20020102531A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 583
SOFTWARE: PatentIn version 3.0
SEQ ID NO 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-964-824A-101
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US-09-964-824A-563
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                   IITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
IITLE OF INVENTION: Sets
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                                                  TITLE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SEQ ID NO 563
LENGTHARE: PATENTIN VERSION 3.0
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APPLICANT: Horrigan, Stephen
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; ORGANISM: Homo sapiens
US-09-964-824A-563
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                                                                                                                         APPLICANT: HOCKLEY, Joseph G.
APPLICANT: Wockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TILLS OF INVENTION: Gene Expression Profiles in Liver Cancer:
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR PILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE PARENTIN VET: 2.1
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US-09-880-107-3420
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2.8%; Score 215.6; DB 10; Length
Best Local Similarity 65.6%; Pred. No. 4.4e-54;
Matches 449; Conservative 0; Mismatches 204; Indels
                              ; Sequence 3420, Application US/09880107; Patent No. US20020142981A1; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
US-09-880-107-3420
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6999 AGGTATTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
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1656 CACCTCCCTGCTCAGTGCTTGGGCTCCACGGGGGGGGTCAGAGCACTCCCTAATTTATG 1715
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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APPLICANT: Lodes, Michael J.
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APPLICANT: Becrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Yngiu
APPLICANT: Smith, Carole Lynn
APPLICANT: Ming, Gordon E.
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APPLICANT: Clapper, Jonathan D.
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
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CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
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Patent No. US20020076414A1
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; ORGANISM: Homo sapiens
US-09-922-217-1105
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| Sequence 192, Application US/09967768A
| Patent No. US20030150877A1
| Patent No. US20030150877A1
| Patent No. US20030150877A1
| APPLICAMTON: Address Meena
| ITLE OF INVENTION: Sets
| FILE REFERENCE: 689280-0.28
| FILE REPERENCE: 689280-0.28
| FILE REPERENCE: 689280-0.28
| PRIOR PELICATION NUMBER: US/60/236, 109
| PRIOR PLILING DATE: 2000-09-28
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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ORGANISM: Homo sapiens
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7398 ATGAGACCCACÇTGCAGATGCCTGGCCTCAGCCAAGGAGGCTGGGGAGACTGTGGCAGGA 7457
                                        7458 GACTGCAGGGACGGAGGGGACAGGGTTGTGTCCTCGG----TACTTCCTGGACTGCCTTC 7513
                                                                                                                                                                                                     1598 GAGCACCGTGATGGAGGAGGACAGGGGCTCCAGCACCTTCTTCTGGACTGGCGTT 1657
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: PREESEQ for Windows Version 4.0
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Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Clapper, Jonathan D.
Skeiky, Yasir A. W.
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Smith, Carole L.
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; ORGANISM: Homo sapiens
US-10-025-380-1105
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31;
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA137P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR PRING DATE: 2000-06-07
PRIOR PLING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2010-06-07
PRIOR FILING DATE: 3.1
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: Sequence 756, Application US/10264049
: Publication No. US20040005579A1
; GENERAL INFORMATION:
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6999 AGGTATTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
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PRIOR FILING DATE: 1999-03-19
WINDER OF SEQ ID NOS: 202
SOFTWARE: PATENTIN VEr. 2.1
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APPLICANT: HINZWANN, BERND
APPLICANT: SCHNITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
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CORGANISM: Homo sapiens
US-10-131-410-64
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LENGTH: 2269
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US-10-131-410-64
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                            1579 ACACCTTCCTCCTCCAGACACCTGGACTGAGCCCAAGGAGGCCTGGG--GAGGCCCTAGGG 1636
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7179 GGACAGGCAGGCCTGACGCCCCCTTAACATGGATGTGTTCCCTGTGTTGCTGTAGAGAG 7238
                                                                                    7239 GAAGAACCIGIIGGGCGIGCCCICIGC -- AGICICCICAAGIGCAGCCIIIGGCCIC - 7293
                                                                                                                               1399 AAGCTGATGTTTTGGTGTATTGTCAGCCATCGTCCTGGGACTCGGAGACTATGGCCTCGC 1458
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TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PAION
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
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PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
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Patent No. US20020052308A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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1559 ACACCTTCCTCCTCCAGACACCTGGACTGAGCCAAGGAGGCCTGGG--GAGGCCCTAGGG 1616
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Best Local Similarity 65.6%; Pred. No. 4.9e-54;
Matches 449; Conservative 0; Mismatches 204; Indels 31; Gaps
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APPLICANT: ROSENTHAL, ANDRE
ATTILE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMORS
FILE REFERENCE: SCH-1763
CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT PILING DATE: 2002-04-25
PRIOR APPLICATION NUMBER: 03/466,673
PRIOR FILING DATE: 2000-09-20
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0; Gaps

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6181 TTGGGATAGGAGAGGCCAGGTCGCCTAGAGGAGAGGGCCACCCCAGACTGATAACTGAGG 6240
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Publication No. US20030143668A1

GENERAL INFORMATION:
APPLICANT: SYNCA, MAKIKO
APPLICANT: AKIYAMA, YUTAKA
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OTHER INFORMATION: a, t, c, g, unknown or other
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LOCATION: (1145). (1145)
OTHER INFORMATION: a, t, c, g, unknown or other
US-10-22-798-1601
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OTHER INFORMATION: a, t, c, g, unknown or other
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NAME/KEY: modified_base
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LOCATION: (201)..(1235)
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ORGANISM: Homo sapiens
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                                                                                                                                 NAME/KEY:
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Publication No. US2030235833A1
Sequence 1601, Application US/10292798
Publication No. US2030235833A1
SEGUENCE INFORMATION:
APPLICANT: SURA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: ALWARA, TUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
FILE REPERENCE: 084335/166
CURRENT APPLICATION NUMBER: US/10/292,798
CURRENT FILING DATE: 2002-11-13
FRIOR APPLICATION NUMBER: 10/017,161
FRIOR FILING DATE: 2001-12-18
FRIOR FILING DATE: 2001-12-18
FRIOR FILING DATE: 2001-12-18
FRIOR FILING DATE: 2001-106-18
FRIOR FILING DATE: 2001-106-18
FRIOR FILING DATE: 2010-106-18
FRIOR FILING DATE: 2010-106-18
FRIOR FILING DATE: 2010-106-18
FRIOR FILING DATE: 2010-106-18
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1059 TTTGGCAAGAACTCTAGTGGCTGGAAGGAAGAAGAGGTTGGAGAGGTGGGAATTAAGGA
                                                                                                                                                                                                                                                                                         7119 TCGGGGCTGGACCCCAGGACCTCAGGCATGAACTCCAGAACTGAAGCCTTCCTGGAA
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ORGANISM: Homo sapiens
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LENGTH: 1435
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4267 CCGCTGCAACATGGACGGAGCCACCTTGCAGCTGTGCGCTGGAGGAGCTGCGGCTAGT 4326
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: LONGOUNDS FOR IMMUNOTHERAPY AND
FILLE OF INVENTION: LONGOUNDS FOR IMMUNOTHERAPY AND
FILLE OF INVENTION: LONGOUNDS FOR IMMUNOTHERAPY AND
FILLE OF INVENTION: 210121.4710.12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILLING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                   4327 CTTTGGACCTCTGGGAGACCAGCTCCATGCCCAGCTTCGGGACCTCAGT 4375
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2.3%; Score 176.2; DB 10; Length 626;
Best Local Similarity 85.6%; Pred. No. 2.3e-42;
Matches 196; Conservative 0; Mismatches 33; Indels 0;
                                                                                                    442 cririceccircreegeaaccaacrecareccaecrecaadaccrear 394
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                                                                                                                                                                                                                                    Sequence 853, Application US/09833263
Patent No. US20020110547A1
GENERAL INFORMATION:
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Stolk, John A.
Wang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
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Smith, Carole L.
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Wang, Aijun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
CRGANISM: Homo sapien
US-09-833-263-853
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US-10-025-380-853/c
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LENGTH: 626
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                                                                                                    Query Match
2.4%; Score 185.8; DB 13; Length 1435;
Best Local Similarity 79.4%; Pred. No. 4.7e-45;
Matches 220; Conservative. 0; Mismatches 57; Indels 0;
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85.6%; Pred. No. 2.3e-42;
tive 0; Mismatches 33; Indels 0
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
PILE REPERBERCE: 210121, 471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
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; LOCATION: (1145)

; OTHER INFORMATION: a, t, c, g, unknown or other

US-10-017-161-1953
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Masqher: Magherine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Smith, Carole Lynn
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Wang, Aijun
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US-09-922-217-853
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4267 CCGCTGCAACATGGACGAGCCACCTCTGCAGCTGTGCGCTGGAGGAGCTGCGGCTAGT 4326
                                                                                                                                                                                                                                                                                                                                                                                     622 TACAGAGAAGGCCAGCTGGTTGGGGAACAGCCCCAGTTCTGGTCGAAGACGCAGGTTCT 563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  502 ACGATGTGACATGGATGGCGCCACCCTTGGAATTGTGCCCTTGAGGAGCTGCGTCTGGT 443
                                                                                                                                                                                                                                                                                                                  0; Gaps
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                                                                                                                                                                                                                                                              Query Match 2.3%; Score 176.2; DB 14; Length 626; Best Local Similarity 85.6%; Pred. No. 2.3e-42; Matches 196; Conservative 0; Mismatches 33; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            442 CTTTGGGCCTCTGGGGACCAACTCCATGCCCAGCTGCGAGACCTCACT 394
; FILE REFERENCE: 210121.471C14
; CURRENT APPLICATION WUMBER: US/10/025,380
; CURRENT FILING DATE: 2001-12-19
; NUMBER OF SEQ ID NOS: 1129
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 853
LENGTH: 626
; TYPE: DNA
; ORGANISM: HOMO Sapiens
US-10-025-380-853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Search completed: February 13, 2004, 11:48:40 Job time : 2738.91 secs
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(without alignments)
10448.744 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_MEW_PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
7: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
11: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
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14: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
15: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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11: /cgn2_6/ptodata/1/pubpna/US10A_PUB.SeQ:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2449703 segs, 1841816367 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                   OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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21
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

			App	[00]	AD	ď	App	App	App	App	App	ppi	App	App	Ϋ́	App	Ap
			927.	32.	2290,		282.	282	282	282	282	74, A	101,	563,	3420.	192.	
		Description	Sequence 927	Seguence	Sequence 2290	Sequence 282,	Sequence 282	Seguence	Seguence	Sequence	Seguence 282,	Sequence	Sequence	Seguence	Seguence	Seguence	Sequence 1105
		QI	US-10-305-720-927	US-09-998-598-32	US-09-998-598-2290	US-09-604-287A-282	US-09-339-338-282	US-09-551-621-282	US-10-124-805-282	US-10-007-805-282	US-10-076-622-282	US-10-097-340-74	US-09-964-824A-101	US-09-964-824A-563	US-09-880-107-3420	US-09-967-768A-192	US-09-922-217-1105
		DB	12	10	10	σ	10	11	13	14	15	15	10	10	10	10	σ
	Query	Length	237	451	499	502	502	502	502	502	502	1907	1915	1915	1915	1915	1917
æ	Query	Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
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Seguence 1105, Ap	Sequence 756, App	Sequence 207, App	Sequence 64, Appl	326		Sequence 1953, Ap	ų	ന	14					Sequence 25, Appl	599	1471,	87,	7	10,	7139	4	Sequence 7031, Ap	46, 7	207	Segmence 350, App	ý	b	65	Sequence 65, Appl
US-10-025-380-1105	US-10-264-049-756	US-09-925-301-207	US-10-131-410-64	US-10-060-036-3261	US-10-292-798-1601	US-10-017-161-1953	5 US-10-156-761-1	US-10-291-172-388	US-10-027-632-149050	US-10-027-632-149050	US-10-156-761-5632	US-10-027-632-109965	US-10-027-632-109965	US-09-919-497-25	US-10-388-934-599	US-09-764-868-1471	US-10-175-523-87	S US-10-156-761-1	US-10-187-267A-10		_	US-1	1	US-10-388-263-201	US-10-044-090-350	US-10-439-249-6	US-09-866-034-6	US-10-210-951-65	US-10-211-884-65
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1917	1956		2269		٦		9025608	619		733	1851	2094	2094	2212	2450	10322	130427	9025608	366	402	470	624	1639	2266	2939	3121	3121	3121	3121
100.0	100.0	100.0	100.0	90.5	90.5	90.5	80.0	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	77.1	77.1	77.1	77.1	77.1	77.1	77.1	77.1	77.1	77.1	77.1
21	21	21	21	19	19	19	16.8	16.4	16.4	16.4	16.4	16.4	16.4	16.4	16.4	16.4	16.4	16.4	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2
c 16	c 17	c 18	c 19	c 20	21	22	23	N	c 25	c 26	27	28	50	c 30	31	32	c 33	c 34	35	36	37	G 38	39	c 40	41	42	43	44	45

ALIGNMENTS

RESULT 1 US-10-305-720-927/c

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Sequence 927, Application US/10305720

Publication No. US20040010136A1

GENERAL IMPORMATION:

APPLICANT: Au-Young, Janice K.; Seilhamer, Jeffrey J.

TITLE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Expressing TILE OF INVENTION: COMPOSITION OF THE REPRENCE: 2002-11-26

PRIOR PLING DATE: 1998-01-30

CURRENT FILING DATE: 1998-01-30

NUMBER OF SEQ ID NOS: 1490

SOFTWARE: PERL Program

SOFTWARE: PERL Program

SOFTWARE: PERL Program

SOFTWARE: PERL Program

SOFTWARE: PERL Program

SOFTWARE: POST

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RESULT 6
US-09-551-621-282/c
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US-09-339-338-282/c
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                                                                                       TYPE: DNA
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Sugan L.
APPLICANT: Harlocker, Sugan L.
APPLICANT: Harlocker, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470C7
CURRENT APPLICATION NUMBER: US/09/604,287A
CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
APPLICANT: Xu, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
FITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REPERENCE: 210121.561
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT APPLICATION NUMBER: US/09/998,598
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: Corixa Invention Disclosure Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AND METHODS FOR THE THERAPY AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
100.0%; Score 21; DB 10; Length 499;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 21; Conservative 0; Mismatches 0; Indels C
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100.0%; Score 21; DB 10; Length 451;
Best Local Similarity 100.0%; Pred. No. 0.73;
Matches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2290, Application US/09998598
Parent No. US200201150922A1
GENERAL INFORMATION:
APPLICANT: Stolk, John A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER FILE REFERENCE: 2.00121.561
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT PILING DATE: 2.001-11-16
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: COCIXA INVENTION Disclosure Database
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21; Conservative
                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
CORGANISM: Homo sapiens
US-09-998-598-32
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; ORGANISM: Homo sapiens
US-09-998-598-2290
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US-09-604-287A-282/c
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US-09-998-598-2290
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LENGTH: 451
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Sequence 282, Application US/0933933A
; Sequence 282, Application US/0933933A
; Patent No. US20020102602A1
; GENREAL INFORMATION:
    APPLICANT: Widiu, Jang
    APPLICANT: Mitcham, Jennifer L.
    APPLICANT: Mitcham, Jennifer L.
    APPLICANT: Mitcham, Jennifer L.
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APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchum, Jennifer L.
APPLICANT: Xu, Jiangchum, Jennifer L.
APPLICANT: Xu, Jiangchum Jennifer L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: COMPOSITIONS FOR THEIR USE
FILE REPERENCE: 210121.470C5
CURRENT APPLICATION NUMBER: US/09/551,621
CURRENT FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 479
SOFTWARE: FastSEQ for Windows Version 3.0
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SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
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Matches 21; Conservative
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ORGANISM: Homo sapiens
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US-09-604-287A-282
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RESULT 9
US-10-076-622-282/c
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APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Fanger, Gary R.
APPLICANT: Penger, Gary R.
APPLICANT: Penger, Tricker B.
APPLICANT: Penger, Gary R.
APPLICANT: Wordil, Patricia D.
APPLICANT: McNeill, Patricia D.
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APPLICANT: McNeill, Patricia D.
APPLICANT: Mc
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; Publication No. U820030166022A1
; Garrard. INFORMATION:
; APPLICANT: Houghton, Raymond L.
; APPLICANT: Bleath, Paul R.
; APPLICANT: Persing, David H.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
; TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
; FILE REFERENCE: 210121.470C12
; CURRENT FILING DATE: 2002-04-15
; NUMBER OF SEQ ID NOS: 627
; SOFTWARE FEASTE FASTER for Windows Version 4.0
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Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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Publication No. US20020150581A1
GENERAL INFORMATION:
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CRGANISM: Homo sapiens
US-10-007-805-282
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) ORGANISM: Homo sapiens
US-10-124-805-282
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LENGIH: 502
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LENGTH: 502
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APPLICANT: Karen GLATT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
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Sequence 282, Application US/10076622
; Sequence 282, Application US/10076622
; Publication No. US20030023036A1
; GENERAL INFORMATION:
   APPLICANT: Houghton, Raymond L.
; APPLICANT: Bleath, Paul R.
; APPLICANT: Peresing, David H.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
; FILE REFERENCE: 210121.470C11
; CURRENT APPLICATION NUMBER: US/10/076,622
CURRENT FILING DATE: 2002-02-13
; NUMBER OF SEQ ID NOS: 627
; SEQ ID NO 282
; SEQ ID NO 282
; LENGTH: 502
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CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-26
PRIOR PLING DATE: 2001-03-26
PRIOR PELING DATE: 2001-03-26
PRIOR PELING DATE: 2001-03-14
PRIOR PELING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-19
PRIOR PLING DATE: 2001-09-26
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PRIOR PELING DATE: 2001-09-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    458 GTACCTCATGGCCCGGCTCAG 438
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Robert C. BAST, Jr.
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Rosemarie SCHMANDT
Xumei ZHAO
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Michael MORRISEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
; ORGANISM: Homo sapiens
US-10-076-622-282
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Sequence 1922, Application US/09967768A

Sequence 1922, Application US/09967768A

Parent No. US20020150877A1

GENERAL INPORMATION:

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

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FILE REPERENCE: 689290-72

CURRENT FILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

SOFTWARE: Patentin version 3.0

SEQ ID NO 192

LENGTH: 1915
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Sequence 14/0.422002014281A1

Patent No. US2002014281A1

GENERAL INFORMATION:

APPLICANT: Horne, Darci T.

APPLICANT: Scherf, Uwe

APPLICANT: Scherf, Uwe

APPLICANT: Gene Logic, Inc.

TILE OF INVENTION: Gene Expression Profiles in Liver Cancer

FILE REFERENCE: 44921-5028-WO

CURRENT FILING DATE: 2010-06-14

PRIOR APPLICATION NUMBER: US 60/211,379

PRIOR PILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

SOFTWARE OF SEQ ID NOS: 3950

SOFTWARE PATENTY NOW NOWER: US 60/237,054

PRIOR FILING DATE: 2000-10-02

NUMBER OF SEQ ID NOS: 3950

SOFTWARE PATENTY NOW NOWER: 2.1
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US-09-880-107-3420
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                      100.0%; Score 21; DB 10; Length 1915; 100.0%; Pred. No. 0.65; 0; Indels 0.
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Matches 21; Conservative
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US-09-967-768A-192
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ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-967-768A-192/c
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US-09-880-107-3420/c
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LENGTH: 1915
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Sequence 5.37, Application US/09964824A

Fatent No. US2002010231A1

GENERAL INFORMATION:

APPLICANT: Horrigan, Stephen

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu

TITLE OF INVENTION: Set 8

TITLE OF INVENTION: Set 8

TITLE OF INVENTION: Set 8

TITLE OF INVENTION WIMBER: US/09/964,824A

CURRENT APPLICATION NUMBER: US/60/236,033

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR FILING DATE: 2000-09-28

SPRIOR FILING DATE: 2000-09-28

SPRIOR FILING DATE: 2000-09-28

SPRIOR FILING DATE: 2000-09-28

SPRIOR FILING DATE: 1000-09-28

SEQ ID NOS: 583

LENGTH: 1915

    Sequence 101, Application US/09964824A
    Patent No. US20020102531A1
    GENERAL INFORMATION:
    TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatulities of INVENTION: Sets

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                                                                                                                                                                      Query Match 1907; Score 21; DB 15; Length 1907; Best Local Similarity 100.0%; Pred. No. 0.65; Matches 21; Conservative 0; Mismatches 0; Indels 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT PILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR STLING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
NUMBER OF SEQ ID NOS: 583
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                                                     TYPE: DNA
CORGANISM: Homo sapiens
US-10-097-340-74
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US-09-964-824A-563
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US-09-964-824A-101
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US-09-964-824A-101/c
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US-09-964-824A-563/c
SEQ ID NO 74
LENGTH: 1907
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LENGTH: 1915
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US-09-021-217-1105/c
Sequence 1105, Application US/09922217
GENERAL INFORMATION:
APPLICANT: Mouse, Michael J.
APPLICANT: Secriter, Heather
APPLICANT: Secriter, Heather
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sitolk, John A.
APPLICANT: Sitolk, John A.
APPLICANT: Sitolk, John A.
APPLICANT: Sitolk, Carole Lynn
APPLICANT: Sitolk, Carole Lynn
APPLICANT: Sitolk, Carole Lynn
APPLICANT: Sitolk, Carole Lynn
APPLICANT: Carole Lynn
APPLICANT: Carole Lynn
APPLICANT: Wang, Ajun
APPLICANT: Wang, Ajun
APPLICANT: Wang, Ajun
APPLICANT: Wang, Ajun
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APPLICANT: Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wang, Wa
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Best Local Similarity 100.0%; Pred. No. 0.65;
Matches 21; Conservative 0; Mismatches 0; Indels C
     0; Mismatches
                                                                                                 Matches 21; Conservative
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ORGANISM: Homo sapiens
US-09-922-217-1105
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Search completed: February 13, 2004, 11:48:34 Job time: 16.4034 secs

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Sequence 282, App
Sequence 282, App
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Sequence 1601, Ap
Sequence 1953, Ap
Sequence 74, App N
Sequence 101, App
Sequence 563, App
Sequence 3420, App
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Sequence 2290, Ap
Sequence 282, App
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10448.744 Million cell updates/sec
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| cgn2 6/ptodata/1/pubpna/PCT PUBCOMB.seq:*
| cgn2 6/ptodata/1/pubpna/PCT NEW PUB.seq:*
| cgn2 6/ptodata/1/pubpna/DCG NEW PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USOG NEW PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USOG NEW PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USOG NEW PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USOB PUBCOMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-98-598-2290

US-09-339-328-2282

US-09-339-338-282

US-09-551-621-282

US-10-124-885-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-097-340-74

US-09-964-824A-101

US-09-984-824A-563

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Maximum Match 100%
Listing first 45 summaries
                                                                                                           OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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21
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Match Length DB
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16 21 100.0 1917 9 US-09-922-217-1105 Sequence 1105, Ap 21 100.0 1917 14 US-10-025-380-1105 Sequence 1105, Ap 22 1 100.0 1956 12 US-10-25-380-1105 Sequence 207, App 22 1 100.0 1956 12 US-10-25-380-1105 Sequence 207, App 22 1 100.0 1956 12 US-10-131-410-64 Sequence 207, App 22 1 100.0 2269 12 US-10-131-410-64 Sequence 207, App 22 1 100.0 2269 12 US-10-131-410-64 Sequence 207, App 22 1 100.0 2269 12 US-10-131-410-64 Sequence 21324, App 22 1 10.8 80.0 792 13 US-10-027-632-138998 Sequence 21322, App 22 15.8 80.0 792 13 US-10-027-632-138998 Sequence 138998, Sequence 138998, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, Sequence 213899, App 23 15.8 80.0 30256 15 US-10-126-761-1 Sequence 21, App1 25 12 US-10-126-2309, Sequence 21, App1 25 12 US-10-156-2309, Sequence 2190, App 23 16.4 78.1 322 12 US-10-156-347 Sequence 2190, App 24 16.4 78.1 352 12 US-10-156-347 Sequence 2190, App 25 16.4 78.1 352 12 US-10-369-493-3351 Sequence 21, App1 26 16.4 78.1 3653 12 US-10-369-493-3351 Sequence 21, App1 26 16.4 78.1 3653 12 US-10-369-493-3351 Sequence 21, App1 26 16.2 77.1 361 9 US-09-38-380 Sequence 2303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 2303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 36 16.2 77.1 361 9 US-09-38-380 Sequence 303, App 26 16.2 77.1 361 9 US-09-38-380 Sequen
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US-10-060-036-3261

US-10-060-036-3261

Sequence 3261, Application US/10060036

Publication No. UG20030031344A1

GENERAL INFORMATION:

APPLICANT: Benson, Darin R.

APPLICANT: Lodes, Michael D.

APPLICANT: Lodes, Michael D.

APPLICANT: Hepler, William T.

APPLICANT: Joing, Yudiu

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

CURRENT FILING DATE: 2002-01-30

NORTHARE: FRACES (10 NOS: 4560

SEQ ID NO 3261

SEQ ID NO 3261

TYPE: DNA

ORGANISM: Homo sapiens

US-10-06-036-3261

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DO CCGGGACATCCTCATCCACCC 21

DA 109 CCGGGACATCCTCATCCACC (129

RESULT 2

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RESULT 2

SEQUENCE 2290, Application US/09998598

SEGENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 315
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US-10-124-805-282
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LENGTH: 502
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Sequence 282, Application US/0933938A

Patent No. US20020102602A1

GENERAL INFORMATION:
APPLICANT: Yudiu, Jiang
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: MITCHAM, DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
CURRENT APPLICATION UNDERS: US/09/339,338A

CURRENT PILING DATE: 1999-06-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Janinifer L.
APPLICANT: Mitcham, Janinifer L.
APPLICANT: Harlocker, Suean L.
APPLICANT: Harlocker, Suean L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
TITLE OF ILLING DATE: 20121.47007
CURRENT APPLICATION NUMBER: US/09/604,287A
CURRENT APPLICATION NUMBER: US/09/604,287A
NUMBER OF SEQ ID NOS: 488
SOFTWARE: FREEER PRESEQ FOR WINDOWS Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
APPLICANT: Stolk, John A.
APPLICANT: Xu, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.561
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT PILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: Corixa Invention Disclosure Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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US-09-604-287A-282
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US-09-339-338-282
                                                                                                                                                                                                                                                                                                                            SEQ ID NO 2290
LENGTH: 499
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LENGTH: 502
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APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.470C5
CURRENT APPLICATION NUMBER: US/09/551,621
CURRENT PILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 4779
SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Score 21; DB 13; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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100.0%; Score 21; DB 11; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels C
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Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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APPLICANT: Sleath, Paul R.
APPLICANT: Sleath, Paul R.
APPLICANT: Steath, Paul R.
APPLICANT: Persing, David H.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470C12
CURRENT APPLICATION NUMBER: US/10/124,805
CURRENT FILING DATE: 2002-04-15
SUFFRARE: FASTSEQ for Windows Version 4.0
SOTTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 282, Application US/10124805; Publication No. US20030166022A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-551-621-282

Sequence 282, Application US/09551621

Publication No. US20030104366A1

GENERAL INFORMATION:

APPLICANT: Yuqui, Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          290 CCGGGACATCCTCATCCACCC 310
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ORGANISM: Homo sapiens
US-10-124-805-282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-551-621-282
                                                                                           TYPE: DNA
CRGANISM: Homo sapiens
US-09-339-338-282
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Sequence 1601, Application US/10292798
| Sequence 1601, Application US/10292798
| Publication No. US2003235833A1
| GENERAL INFORMATION:
| APPLICANT: SUWA, MAKIXO
| APPLICANT: SUWA, TUTAKA
| APPLICANT: ABURATNANI, HIROYUKI
| TITLE OF INVENTIONS GIANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
| FILE REFERENCE: 084335/16
| CURRENT FELING DATE: 2000-11-13
| PRIOR PILING DATE: 2001-12-18
| PRIOR PILING DATE: 2001-12-18
| PRIOR PILING DATE: 2001-06-18
| NUMBER OF SEQ ID NOS: 2070
| SOFTWARE: Patentin Ver. 2.1
| SEQ ID NO 1601
| WARDER OF TABORY FAILST
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Squence 1953, Application US/10017161

Publication No. US203030143668A1

GENERAL INFORMATION:

APPLICANT: SUMA, MAXIKO

APPLICANT: AKIYAMA, YUTAKA

APPLICANT: AKIYAMA, YUTAKA

APPLICANT: AKIYAMA, YUTAKA

APPLICANT: ARINAMA, YUTAKA

APPLICANT: AKIYAMA, YUTAKA

CURRENT APLICATION NOWEL G PROTEIN-COUPLED RECEPTORS

FILE REFERENCE: 084335/0152

CURRENT APPLICATION NUMBER: US/10/017,161

CURRENT APPLICATION NUMBER: US/2001/246789

PRIOR FILING DATE: 2001-06-18

PRIOR FILING DATE: 2001-06-18

NUMBER OF SEQ ID NOS: 2430

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 1953

TENOTFI 1435
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OTHER INFORMATION: a, t, c, g, unknown or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LOCATION: (1145)..(1145); OTHER INFORMATION: a, t, c, g, unknown or other US-10-292-798-1601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                335 CCGGGACATCCTCATCC 315
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LOCATION: (201)..(1235)
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FEATURE:
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; Sequence 282, Application US/10076622
; Publication No. US20030023036A1
; GENERAL INFORMATION:
    APPLICANT: Houghton, Raymond L.
    APPLICANT: Houghton, Raymond L.
    APPLICANT: Pershing, David H.
    ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
    ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
    ITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
    ITLE REPERBNCE: 210121.47011
    CURRENT FILING DATE: 2002-02-13
    NUMBER OF SEG ID NOS: 627
    SOFTWARE: FastSEQ for Windows Version 4.0
    SEQ ID NO 282
    LENGTH: 502
                                                                                                                                                                                                                                                                  APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Marcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Henderson, Robert A.
APPLICANT: Panger, Gary R.
APPLICANT: Powdick, Thomas S.
APPLICANT: Wordill, Patricia D.
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: WORGILL.470Cl0
CURRENT FILIE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: COMPOSITIONS AND MAGNOSI
CURRENT FILIE STOLE SOLI - 2-07
NUMBER OF SEQ ID NOS: 593
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 502
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100.0%; Score 21; DB 14; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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                                                   290 ccecedacarcercarceacec 310
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1 CCGGGACATCCTCATCCACCC 21
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; ORGANISM: Homo sapiens
US-10-076-622-282
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CRGANISM: Homo sapiens
US-10-007-805-282
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US-09-964-824A-563
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 21; DB 13; Length 1435; Best Local Similarity 100.0%; Pred. No. 2.4; Matches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                NAME/KEY: modified base
LOCATION: (1040)...(1139)
OTHER INFORMATION: a, t, c, g, unknown or other
                                                                                                                                                                                                                                                                             LOCATION: (1145)
COTHER INFORMATION: a, t, c, g, unknown or other US-10-017-161-1953
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PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-14
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-16
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PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DATE: 2001-09-17
PRIOR FILING DATE: 2001-09-18
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
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Publication No. US20030087250A1
GENERAL INFORMATION:
APPLICANT: John MONHAN
APPLICANT: Shobstian HOERSCH
APPLICANT: Shubhangi KAWATKAR
APPLICANT: Shubhangi KAWATKAR
APPLICANT: Steve G. KOVATS
APPLICANT: Machel MORRISEY
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
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Robert C. BAST, Jr.
Karen LU
Rosemarie SCHMANDT
Xumei ZHAO
                                                                                                                                                                                                                                                        NAME/KEY: modified_base
                                                                                LOCATION: (201)..(1235)
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ORGANISM: Homo sapiens
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LOCATION: (1)..(1435)
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Best Local Similarity
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                          FEATURE:
NAME/KEY:
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APPLICANT: Horrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat-
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
FILLE REPERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 21; DB 10; Length 1915; 100.0%; Pred. No. 2.3;
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0; Indels
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0; Mismatches
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                                                                 932 CCGGGACATCCTCATCCACCC 952
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                                 1 CCGGGACATCCTCATCCACCC 21
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SEQ ID NO 563
LENGTH: 1915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21; Conservative
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US-09-964-824A-563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: DNA; ORGANISM: Homo sapiens
US-09-964-824A-101
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Best Local Similarity
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Sequence 192, Application US/09967768A

Sequence 192, Application US/09967768A

Sequence 192, Application US/09967768A

GENERAL INFORMATION:

APPLICANT: Adjusture, Meens

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using signatu

TITLE OF INVENTION: Sets

FILE REFERENCE: 689290-72

CURRENT APPLICATION NUMBER: US/09/967,768A

CURRENT FILING DATE: 2001-09-28

PRIOR APPLICATION NUMBER: US/60/236,034

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

SOFTHARE: PatentIn Version 3.0

SEQ ID NO 192

LEMOTH: 1915

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                                                                               Sequence 3420, Application US/09880107
Patent No. USZO020142981A1
Patent No. USZO020142981A1
APPLICANT: Workley, Joseph G.
APPLICANT: Workley, Joseph G.
APPLICANT: Scherf, USCALEY, Joseph G.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENITON: Gene Expression Profiles in Liver Cancer:
FILE REFERENCE: 4429-15.028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843
US-09-880-107-3420
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CORGANISM: Homo sapiens
US-09-967-768A-192
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ORGANISM: Homo sapiens
RESULT 14
US-09-880-107-3420
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US-09-967-768A-192
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Search completed: February 13, 2004, 11:48:25

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Publication No. US20040010136A1

Publication No. US20040010136A1

Publication No. US20040010136A1

Publication No. US20040010136A1

APPLICANT: Au-Young, Janice K.; Seilhamer, Jeffrey J.

TITLE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Expressic TITLE OF INVENTION: COMPOSITION OF THE REFERENCE: PA-0002-1 CON; CURRENT APPLICATION NUMBER: US/10/305,720

CURRENT FILING DATE: 2002-11-26

PRIOR FILING DATE: 1998-01-30

NUMBER OF SEQ ID NOS: 1490

SEQ ID NOS: 1490

SEQ ID NO 927

LENGTH: 237
                                   Sequence 32, Appl
Sequence 2290, Ap
Sequence 2373, Ap
Sequence 2379, Ap
Sequence 74, Appl
Sequence 101, App
Sequence 563, App
Sequence 563, App
Sequence 192, App
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Sequence 64, Appl
Sequence 40, Appl
Sequence 1730, Ap
Sequence 3052, Ap
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Sequence 678, App
Sequence 33421, A
Sequence 3, Appli
Sequence 125191,
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Sequence 1105, Ap
Sequence 756, App
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Sequence 760, App
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Sequence 167020,
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Sequence 4493, Ap
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Sequence 13
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; OTHER INFORMATION: Incyte ID No. US20040010136A1 773734
US-10-305-720-927
12 US-10-305-720-927
15 US-09-988-598-32
15 US-09-988-598-32
16 US-09-988-52290
17 US-09-964-8248-1563
18 US-10-066-543-3239
18 US-09-964-8248-163
10 US-09-964-8248-163
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              US-10-305-720-927
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            February 12, 2004, 21:34:51; Search time 25.2461 Seconds (without alignments) 2334.540 Million cell updates/sec
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1. /cgn2_6/ptodata/1/pubpna/PCT_NWA PUB.seq:*
2. /cgn2_6/ptodata/1/pubpna/PCT_NWA PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/USO6_NWW PUB.seq:*
4. /cgn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
5. /cgn2_6/ptodata/1/pubpna/USO6_NWW PUB.seq:*
6. /cgn2_6/ptodata/1/pubpna/USO8_NWW PUB.seq:*
7. /cgn2_6/ptodata/1/pubpna/USO8_NWW PUB.seq:*
8. /cgn2_6/ptodata/1/pubpna/USO8_NWW PUB.seq:*
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                                           GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                     OM protein - nucleic search, using frame_plus_p2n model
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Maximum Match 100%
Listing first 45 summaries
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Kgapop 10.0 , Kgapext
Kgapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 2000000000
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84
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Match Length DB
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                                                                                                                                                                                                                                                                                                                                                Perfect score:
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                                                                                                                                                                                                                                                                                                                         Title:
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Page

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US-09-998-588-2290/C

Sequence 2290, Application US/09998598

Patent No. US20020150922A1

GENERAL INFORMATION:
APPLICANT: Stolk, John A.
APPLICANT: AL, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
CHENERAL OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.561

CURRENT PILION DATE: 2001-11-16

NUMBER OF SEQ ID NOS: 2606

SOFTWARE: COCINX INVENTION DISCLOSURE DATABASE

COTRACTH: 499
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Xu, Jiangchun

APPLICANT: Chenault, Madelein Joy

APPLICANT: Meagner, Madelein Joy

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER

FILE REFERENCE: 210121.561

CURRENT APPLICATION VUMBER: US/09/998,598

CURRENT PILING DATE: 2001-11-16

NUMBER OF SEQ ID NOS: 2606

SOUTHARRE: Corixa Invention Disclosure Database

SEQ ID NO 32

LENGTH: 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            170 AAAAACTCAAGCGGCTGGAAGAGGAAGAAGTTCTCCAGAGTCGGAAC 217
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Matches:
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Mismatches:
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      Length:
Matches:
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Gaps:
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Patent No. US20020150922A1
GENERAL INFORMATION:
3.23e-06
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CORGANISM: Homo sapiens
US-09-998-598-2290
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ORGANISM: Homo sapiens
US-09-998-598-32
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Best Local Similarity:
Query Match:
                                           Percent Similarity:
Best Local Similarity:
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Pred. No.:
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Pred. No.:
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DB:
      Pred. No.:
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Sequence 2379, Application US/10060036

Publication No. US20030073144A1

SERBERLI INFORMATION:
APPLICANT: Benson, Darin R.
APPLICANT: Persing, Michael D.
APPLICANT: Persing, David H.
APPLICANT: Hepler, William T.
APPLICANT: Obvid H.
APPLICANT: Transp. Yuqiu
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER
FILE REFERENCE: 210121.566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 63 AAAAACTCAAGCGGCTGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 110
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   Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
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                                                                                                    US-08-978-217-12 (1-16) x US-09-998-598-2290 (1-499)
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: 438
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                    APPLICANT: Jiang, Yuqiu
APPLICANT: Pyle, Ruth A.
APPLICANT: Xu, Jiangchun
APPLICANT: Indirias, Carol Yoseph
APPLICANT: Lodes, Michael J.
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Percent Similarity: 100.00% Best Local Similarity: 100.00% Query Match: 100.00% DB:
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Best Local Similarity:
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US-09-964-824A-101
US-09-964-824A-101
US-09-964-824A-101
Sequence 101, Application US/09964824A
Sequence 101, Application US/09964824A
Sequence 101, Application US20020102531A1
Sequence 102.0020102531A1
Sequence 102.0020102531A1
SEQUENCE INFORMATION:
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR PELLING DATE: 2000-09-28
PRIOR PELLING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR PELLING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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Patent No. US20020102531A1

Ratent No. US20020102531A1

RATENERAL INFORMATION:
APPLICANT: HORITION:
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturities of INVENTION: Sets

TITLE OF INVENTION: AS ets
FILE REFERENCE: 69290-73

CURRENT APPLICATION NUMBER: US/69/964,824A

CURRENT PILING DATE: 2001-09-27

PRIOR RILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28
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Mismatches:
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Matches:
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100.00%
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   ; ORGANISM: Homo sapiens
JS-10-097-340-74
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                                                                                                                                                               Percent Similarity:
Best Local Similarity:
Query Match:
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TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
CURRENT APPLICATION NUMBER: G0/276,025
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR PELING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR PRIOR APPLICATION NUMBER: 60/311,732
PRIOR PRIOR PRIOR DATE: 2001-09-26
PRIOR PRIOR DATE: 2001-09-26
PRIOR PRIOR PRIOR DATE: 2001-09-26
PRIOR PRIOR DATE: 2001-09-26
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PRIOR PRIOR PRIOR DATE: 2001-09-26
PRIOR PRIOR PRIOR DATE: 2001-09-26
PRIOR PRIOR PRIOR DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
PROPERTY NOTARE PRESEQ FOR Windows Version 4.0
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Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION NUMBER: US/10/060,036 CURRENT FILING DATE: 2002-01-30 NUMBER OF SEQ ID NOS: 4560 SOFTWARE: PastSEQ for Windows Version 4.0 SSOFTWARE: 620
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 74, Application US/10097340 Publication No. US20030087250A1 GENERAL INFORMATION: APPLICANT: John MONAHAN
                                                                                                                                                                                                                                                   NAME/KEY: misc_feature

) ICCATION: 169, 456, 487, 488, 602

; OTHER INFORMATION: n = A,T,C or G

US-10-060-036-2379
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Sebastian HOERSCH
Shubhangi KAWATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gordon B. MILLS
Robert C. BAST, Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosemarie SCHMANDT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Xumei ZHAO
Karen GLATT
                                                                                                                                                                                                              ORGANISM: Homo sapiens
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Best Local Similarity:
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LENGTH: 1907
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DB:
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1187 AAAAACTCAAGGGCTGGAAGGAGGAAGAAGCTTCTCCCAGAGTCGGAAC 1234
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
PILE REPERENCE: 210121, 471C13
CURRENT APPLICATION NUMBER: US/09/922, 217
CURRENT FILING DATE: 2001-08-03
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  CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,109
PRIOR PILING DATE: 2000-09-28
PRIOR PRIOR TILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER: PRIOR PILING DATE: 2000-09-28
SOFWARE: PALCATION NUMBER: US/60/236,111
SEQ ID NO 192
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps:
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SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. US20020076414A1
GENERAL INFORMATION:
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Stolk, John A.
Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Homo sapiens
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Best Local Similarity:
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US-09-922-217-1105
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                                                                                                                                                                                                                                                               US-09-967-768A-192
                                                                                                                                                                                                      LENGTH: 1915
                                                                                                                                                                                                                                                                                                     Alignment Scores:
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LENGTH: 1917
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DB:
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No.
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FILLE REFERENCE: 689290-72
CURRENT APPLICATION NUMBER: US/09/967,768A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3420, Application US/09880107
Patent No. USZO020142981A1
Patent No. USZO020142981A1
APPLICANT: Horne, Darci T.
APPLICANT: Workley, Joseph G.
APPLICANT: Scherf U. Joseph G.
APPLICANT: Scherf U. Joseph G.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer: FILE REFERENCE: 4421-5228-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PATENTIN VONE: 2.1
SEQ ID NO 3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1185 AAAAACTCAAGCGGCTGGAAGGAGGAAGAGGGTTCTCCCAGAGTCGGAAC 1232
                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843 US-09-880-107-3420
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US-09-967-768A-192
Sequence 192, Application US/09967768A
Patent No. US20020150877A1
GENERAL INFORMATION:
APPLICANT: AUGUSTUS.
APPLICANT: AUGUSTUS.
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
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ORGANISM: Homo sapiens
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Best Local Similarity:
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                                        SEQ ID NO 563
LENGTH: 1915
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Conservative:

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Percent Similarity:
Best Local Similarity:
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LENGTH: 2269
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US-10-131-
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                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Vedvick Thomas S.
APPLICANT: Carrer, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121-471C14
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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Publication No. US20040005579A1
FUBLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PAI37P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR FILING DATE: 2001-06-07
PRIOR FILING DATE: 2001-06-07
PRIOR FILING DATE: 2001-06-07
PRIOR SEDIECTION NUMBER: US 60/209,467
PRIOR SEDIECTION NUMBER: US 60/209,467
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
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                                   Sequence 1105, Application US/10025380
Publication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Loddes, Michael J.
APPLICANT: Genes, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Reagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                 Clapper, Jonathan D.
Skeiky, Yasir A. W.
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Vedvick Thomas S.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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CNGANISM: Homo sapiens
US-10-264-049-756
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Best Local Similarity:
Query Match:
                    -10-025-380-1105
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Pred. No.:
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LENGTH: 1956
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APPLICANT: STEAM'S BERND
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PLIARSKY, CHRISTIAN
APPLICANT: PLIARSKY, CHRISTIAN
APPLICANT: PLIARSKY, CHRISTIAN
APPLICANT: DAHL, EDGAR
APPLICANT: ROSENTHAL, ANDRE
TITLE OP INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUB OF BREAST
TITLE OP INVENTION: TUMORS
FILE REFERENCE: SCH-1763
CURRENT PILLING DATE: 2002-04-25
CURRENT PILLING DATE: 2000-09-20
PRIOR PILLING DATE: 2000-09-20
PRIOR PILLING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 202
SUFFRIENCE SEQ ID NOS: 202
SUFFRIENCE PATCHILING VET: 2.1
                                                                                                                                                                                                                      1226 AAAAACTCAAGCGGCTGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 1273
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                                                                                                                                                                    1 LysAsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                                                                                                                                                                                                                                     Sequence 207, Application US/09925301

Sequence 207, Application US/09925301

Sequence 207, Application US/09925301

Patent No. US2002062308A1

GENERAL INFORMATION:

TILE OF INVENTION: Nucleic Acids, Proteins and Antibodies

FILE REFERENCE: PA106

CURRENT APPLICATION NUMBER: US/09/925,301

CURRENT PILING DATE: 2001-08-10

PRIOR FILING DATE: 2000-03-08

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1694

SEQ ID NO 207

LENGTH: 1996
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Matches:
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Mismatches:
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                          Mismatches:
Indels:
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; ORGANISM: Homo sapiens
US-09-925-301-207
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ORGANISM: Homo sapiens
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US-10-131-410-64

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	Length:	Matches:	Conservative:	Mismatches:	Indels:	Gaps:
	3.63e-05	84.00	100.00\$	100.00\$	100.00%	12
Alignment Scores:	Pred. No.:	Score:	Percent Similarity:	Best Local Similarity:	Query Match:	DB:

US-08-978-217-12 (1-16) x US-10-131-410-64 (1-2269)

Search completed: February 13, 2004, 01:46:08 Job time : 28.2461 secs

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Sequence 944, App Sequence 944, App Sequence 944, App Sequence 853, App Sequence 813, App Sequence 101, App Sequence 101, App Sequence 101, App Sequence 102, App Sequence 105, App Sequence 105, App Sequence 105, App Sequence 1016, App Sequence 1016, App Sequence 1019, App Sequence 10194, Sequence 10194, Sequence 10195, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10198, Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App S
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Sequence 139, App
Sequence 139, App
Sequence 139, App
Sequence 92, App
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APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Wang, Aljun
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09.03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FASTSEQ for Windows Version 4.0
                     ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 944, Application US/09922217
Patent No. US20020076414A1
GENERAL INFORMATION:
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Magher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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  US-09-922-217-944/C
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-MODEL=frame+ p2n.model -DEV=xlh
-Q=Cgn2_1/USF0=Cpool/USC08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
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-Q=/Cgn2_1/USF0=Cpool/USC08978217/runat_1002504_1-MATRIX=Dlosum62
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=Dlosum62
-TRAMS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
-THR MAXIEN=0 -ALIGN=15 -MODE=LOOTL -OUTPWF=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=200000000 -USE=USOB978217 @CGN 1 1.690 @runat 10022004 133827_20514
-NCPU=6 -LOOPC=3 -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WALT -DSPBLOCK=100
-LONGLOG -DEV_TIMEOTF=120 -WARN TIMEOTF=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                   February 12, 2004, 21:34:51; Search time 132.542 Seconds (without alignments) 2334.540 Million cell updates/sec
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1 NCALEELRLVFGFLGDQLHA......ELLDDGQQASPYHFGSCGAG
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB_seq:*
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7: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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                          GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                      OM protein - nucleic search, using frame_plus_p2n model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2449703 seqs, 1841816367 residues
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Listing first 45 summaries
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Kgapop 10.0 , Kgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Match Length DB
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Sequence:
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No.
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412 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-978-217-7 (1-84) x US-10-025-380-944 (1-563)
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                                                                                                                                                                                                                                                                                              ; Sequence 944, Application US/10025380; Publication No. US20020182191A1; GENERAL INFORMATION:
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US-10-025-380-944
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Best Local Similarity:
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US-10-025-380-944/c
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Pred. No.:
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LENGTH: 563
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US-09-833-263-944/C

US-09-833-263-944/C

Sequence 944, Application US/09833263

Patent No. US20020110547A1

GENERAL INFORMATION:

APPLICANT: Wang, Aijun

APPLICANT: Clapper, Jonathan D.

APPLICANT: Meagher, Madelaine J.

APPLICANT: Stolk, John A.

APPLICANT: OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

FILE REPERENCE: 210121.471C12

CURRENT APPLICATION UNMER: US/09/833,263

CURRENT FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: PSetSEQ for Windows Version 3.0

SEQ ID NO 944

LENGTH: 563
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Matches:
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Matches:
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Score: 445.00
Percent Similarity: 100.00$
Suber Match: 100.00$
Dery Match: 100.00$
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Percent Similarity: 100.00$
Best Local Similarity: 100.00$
Query Match: 100.00$
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 ; SEQ ID NO 944
; LENGTH: 563
; TYPE: DNA
; CRGANIEM: Homo sapiens
US-09-922-217-944
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CRGANISM: Homo sapien
US-09-833-263-944
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352 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGC 293
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41 GlulysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer
                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Cacrist, Heather
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Mangy Tongtong
APPLICANT: Wangy Tongtong
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Ship, Ajun
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Carter, Darrick
TILLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND SEPLICANT: CHARLEY PRILICAND NUMBER: US/10/025,380
CURRENT PILLING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOUTHARE FEREBEC for Windows Version 4.0
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          APPLICANT: Meagher, Validation J. Madeleine J. TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 21012.17C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 853
LENGTH: 626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 GlnLeuArgAspLeuThrSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeu
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CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
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Matches:
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Publication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodges, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Scotist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Wang, Tongtong
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary K.
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    APPLICANT: Stolk, John A.
                                                                                                                                                                                                                         TYPE: DNA
CRGANISM: Homo sapien
US-09-833-263-853
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Best Local Similarity:
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
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CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
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Patent No. US20020110547A1
GENERAL INFORMATION:
APPLICANT: Wang, Aljun
APPLICANT: Clapper, Jonathan D.
                                                                                                                                                                                                                   APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Beerist, Heather APPLICANT: Benson, Darin R. APPLICANT: Meagher, Madeleine Joy APPLICANT: Stolk, John A. APPLICANT: Wang, Tongtong
                                                                                                                                                            Sequence 853, Application US/09922217
Patent No. US20020076414A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   Jiang, Yuqiu
Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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                                         81 CysGlyAlaGly 84
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CRGANISM: Homo sapiens
US-09-922-217-853
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Best Local Similarity:
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APPLICANT:
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APPLICANT: Rachel E. MEYERS
APPLICANT: Rachel E. MEYERS
APPLICANT: Rachel E. MEYERS
APPLICANT: Peter OLANDT
APPLICANT: Ami SEN
APPLICANT: Ami SEN
APPLICANT: Ami SEN
APPLICANT: Rosemarie SCHWANDT
APPLICANT: Rosemarie SCHWANDT
APPLICANT: Rosemarie SCHWANDT
APPLICANT: Karen LU
APPLICANT: Karen LU
APPLICANT: Karen GLATT
ITILE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
ITILE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
ITILE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
ITILE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
ITILE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
CURRENT FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR PLILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-44
PRIOR FILING DATE: 2001-03-49
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
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Mismatches:
Indels:
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: John MONAHAN
APPLICANT: Marjula GANNAVARAPU
APPLICANT: Sebastian HOERSCH
APPLICANT: Shubhang! KAMATTAR
APPLICANT: Rechel E. MEYERS
APPLICANT: Rachel E. MEYERS
APPLICANT: Peter OLANDT
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                                            TYPE: DNA
CORGANISM: Homo sapiens
US-10-025-380-853
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Best Local Similarity:
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; SEQ ID NO 853
; LENGTH: 626
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Sequence 101, Application US/09964824A

Sequence 101, Application US/09964824A

Patent No. US20020102531A1

GENERAL INFORMATION:
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signation TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signation TITLE OF INVENTION: Sets

FILE REFERENCE: 689290-73

CURRENT APPLICATION NUMBER: US/09/964,824A

CURRENT APPLICATION NUMBER: US/00/236,033

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR FILING DATE: 2000-09-28

NUMBER: US/60/236,028

NUMBER: OF SEQ ID NOS: 583
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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Mismatches:
Indels:
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PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 1907
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SEQ ID NO 101
LENGTH: 1915
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Score:
Bercent Similarity:
Dest Local Similarity:
Query Match:
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ORGANISM: Homo sapiens
                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
US-10-097-340-74
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Sequence 192, Application US/09967768A

Patent No. USZ0020150877A1

GENERAL INFORMATION:
APPLIAL INFORMATION:
APPLIAL INFORMATION:
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-72

CURRENT APPLICATION NUMBER: US/09/967,768A

CURRENT FILING DATE: 2001-09-28
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                                                                                               US-09-880-107-3420

| Sequence 3420, Application US/09880107
| Sequence 3420, Application US/09880107
| Parent No. US20020142981A1
| GENERAL INFORMATION:
| APPLICANT: Horne, Darci T.
| APPLICANT: Scherf, Useph G.
| APPLICANT: Gene Logic, Inc.
| APPLICANT: Gene Logic, Inc.
| TITLE OF INVENTION: Gene Logic, Inc.
| FILE REFERENCE: 4492-5028-WO
| CURRENT APPLICATION NUMBER: US/09/880,107
| FILE REPERINCE: 4301-6-14
| PRIOR PILING DATE: 2000-06-14
| PRIOR PILING DATE: 2000-06-14
| PRIOR PILING DATE: 2000-06-14
| PRIOR PILING DATE: 2000-10-02
| NUMBER OF SEQ ID NOS: 3950
| SEQ ID NO 3420
| LENGTH: 1915
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US-09-880-107-3420
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Conservative:
Mismatches:
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81 CysGlyAlaGly 84
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Best Local Similarity:
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US-09-967-768A-192
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2001-09-28
PRIOR PELING DATE: 2000-09-28
PRIOR PELING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
RIOR APPLICATION NUMBER: US/60/236,028
PRIOR PILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
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Mismatches:
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US-08-978-217-7 (1-84) x US-09-964-824A-101 (1-1915)
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. Sequence 563, Application US/09964824A
; Patent No. US/0020102531A1
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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429 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGGC 488
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APPLICANT: Smith, Carole Lynn
APPLICANT: King, Gordon E.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
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Matches:
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CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
PRIOR APPLICATION NUMBER: US/60/236,109
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,111
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 325
SOFTWARE: Patentin version 3.0
LENGTH: 1915
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Patent No. US20020076414A1
GENERAL INFORMATION:
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Magher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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; ORGANISM: Homo sapiens
US-09-922-217-1105
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; ORGANISM: Homo sapiens
US-09-967-768A-192
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Best Local Similarity:
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431 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGACCAACTCCATGCC 490
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
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SENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodge, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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Best Local Similarity:
Query Match:
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Best Local Similarity:
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US-10-025-380-1105
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Search completed: February 13, 2004, 01:46:05
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PA137P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: US/US01/18569
PRIOR PILING DATE: 2001-06-07
PRIOR PILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
SOFTWARE: Patentin Vos: 4360
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US-08-978-217-7 (1-84) x US-10-025-380-1105 (1-1917)
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Matches:
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; ORGANISM: Homo sapiens
US-10-264-049-756
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US-10-264-049-756
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LENGTH: 1956
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                                                                                                             APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REPREMENCE: PAAG6
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-06
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PAHCHIN Ver. 2.0
SEQ ID NO 207
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                                             ; Sequence 207, Application US/09925301
; Patent No. US20020052308A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
CRGANISM: Homo sapiens
US-09-925-301-207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Percent Similarity:
Best Local Similarity:
RESULT 15
US-09-925-301-207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match:
DB:
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509

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February 13, 2004, 01:27:22 ; Search time 88.8409 Seconds (without alignments) 10448.744 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_TBM* PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/US06_NEW PUB.seq:*
4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
6: /cgn2_6/ptodata/1/pubpna/US08_NEW PUB.seq:*
7: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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11: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
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13: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
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15: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
15: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
16: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
17: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
18: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
18: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2449703 seqs, 1841816367 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                             OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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252
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                              Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Searched:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARIES

	Description	Sequence 944, App	Sequence 944, App	Sequence 944, App	Sequence 853, App	Sequence 853, App	Sequence 853, App	Sequence 74, Appl	Sequence 101, App	Sequence 563, App	Seguence 3420, Ap	Sequence 192, App	Sequence 1105, Ap	Sequence 1105, Ap	Sequence 756, App	Sequence 207, App
Sorrightes	. QI	US-09-922-217-944	US-09-833-263-944	US-10-025-380-944	US-09-922-217-853	US-09-833-263-853	US-10-025-380-853	US-10-097-340-74	US-09-964-824A-101	US-09-964-824A-563	US-09-880-107-3420	US-09-967-768A-192	US-09-922-217-1105	US-10-025-380-1105	US-10-264-049-756	US-09-925-301-207
	DB	6	9	14	σ	10	14	15	10	10	10	10	σ	14	12	σ
	Length DB	563	563	563	626	626	626	1907	1915	1915	1915	1915	1917	1917	1956	1996
o k	Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
	Score	252	252	252	252	252	252	252	252	252	252	252	252	252	252	252
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Query Match 100.0%; Score 252; DB 9; Length 563; Best Local Similarity 100.0%; Pred. No. 2.1e-64; Matches 252; Conservative 0; Mismatches 0; Indels 0

Sequence 64, Appl Sequence 1740, App Sequence 1740, App Sequence 2216, App Sequence 28055, Sequence 280565, Sequence 1356, App Sequence 1356, App Sequence 1356, App Sequence 280564, Sequence 280564, Sequence 280564, Sequence 15959, Sequence 15959, Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App	Sequence 4804, App Sequence 269, App Sequence 1132, App Sequence 1132, App Sequence 1801, App Sequence 2265, App Sequence 145, App Sequence 363, App Sequence 3220, App Sequence 3220, App Sequence 3220, App	AND DIAGNOSIS FOR THEIR USE
US-10-131-410-64 US-09-867-701-4818 US-09-998-598-1740 US-09-998-598-1740 US-10-10-510-356 US-10-10-51-356 US-10-027-632-280565 US-10-027-632-280565 US-10-029-386-11328 US-10-029-386-11328 US-10-029-386-12328 US-10-029-386-12328 US-10-027-632-280564 US-10-027-632-280564 US-10-027-632-280564 US-10-027-632-159559 US-10-027-632-159559 US-10-259-165-233	US-09-233-876-4804 US-09-962-436-269 US-09-954-456-55 US-09-954-456-1132 US-09-954-456-1132 US-09-954-531-145 US-09-954-531-363 US-09-810-808-4 US-09-910-808-4 US-09-100-242-515-3220 US-10-242-515-3220	y IMMUNOTHERAPY CER AND METHODS /922,217 sion 4.0
	70 0000000	US/05/05/05/05/05/05/05/05/05/05/05/05/05/
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	2.000000000000000000000000000000000000	17-944/c 944, Application US/O: USZ0020076414A1 T: Lodes, Michael J. T: Lodes, Michael J. T: Serist, Heather T: Serist, Heather T: Serist, Heather T: Magher, Madleinn T: Stolk, John A. T: Mang, Tongtong T: Yugin T: Stolk, John A. T: Clanger, Vagin T: Wang, Apin T: Clanger, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Clapper, Jonathan T: Filly Gordon B. T: Wang, Aijun T: Clapper, Jonathan T: FastSEQ for Windows 1944 563 M: Homo sapiens M: Homo sapiens
6 / 6 / 6 / 6 / 6 / 6 / 6 / 6 / 6 / 6 /	ჀჅჄჅჅႷႷႷႷႷႷႷ ჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅჅ	1 922 600 - 1 1
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Patent No. US20020110547A1

GENERAL INFORMATION:

APPLICANT Wang, Aijun

APPLICANT: Clapper, John A.

APPLICANT: Clapper, John A.

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

CURRENT APPLICATION NUMBER: US/09/833,263

CURRENT APPLICATION NUMBER: US/09/833,263

CURRENT FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 944

LENGHAL: 563
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                                                                      121 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGG
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Best Local Similarity 100.0
Matches 252, Conservative
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; ORGANISM: Homo sapien
US-09-833-263-944
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US-10-025-380-944/c
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Sequence 944, Application US/10025380
Publication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: Ku, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.

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352 GAGAAGGATGGCATGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210122-471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 252; DB 14; Length 563; Best Local Similarity 100.0%; Pred. No. 2.1e-64; Matches 252; Conservative 0; Mismatches 0; Indels 0
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 853, Application US/0992217
Sequence 853, Application US/0992217
Septent No. US20020076414A1
APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Macher, Madeleine Joy
APPLICANT: Mang, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Jiang, Yuqiu
APPLICANT: Smith, Carole Lynn
APPLICANT: King, Gordon E.
APPLICANT: Wang, Aljun
APPLICANT: Wang, Aljun
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
                                                                         Jiang, Yuqiu
Smith, Carole L.
King, Gordon E.
Wang, Aljun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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US-10-025-380-944
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471 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGGCCTCTGGGGGGACCAACTCCATGCC 412
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CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 853
LENGTH: 626
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                                                                                                                                                                                    100.0%; Score 252; DB 9; Length 626; 100.0%; Pred. No. 2.2e-64; tive 0; Mismatches 0; Indels
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 853, Application US/09833263
; Patent No. US20020110547A1
; GENERAL INFORMATION:
                                                                                                                                                                           Query Match
Best Local Similarity 100.0
Matches 252; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  241 TGTGGCGCAGGA 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         231 TGTGGCGCAGGA 220
                                                                             TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wang, Aijun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , ORGANISM: Homo sapien
US-09-833-263-853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 5
US-09-833-263-853/c
                                                                                                                                US-09-922-217-853
                         SEQ ID NO 853
LENGTH: 626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.41/214
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FASISED for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
                                                                                                                                                                   ; Sequence 853, Application US/10025380; Publication No. US20020182191A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                        APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Masgher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jiang, Yuqiu
APPLICANT: Smith, Carole L.
APPLICANT: King, Gordon B.
APPLICANT: Wang, Ajun
APPLICANT: Sheiky, Xain D.
APPLICANT: Sheiky, Yasir A. W.
APPLICANT: Sheiky, Yasir A. W.
APPLICANT: Clapper, Gordon B.
APPLICANT: Carter Darrie.
241 TGTGGCGCAGGA 252
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Matches 252; Conservative
                                                231 rerececkéck 220
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                                                                                                                                            US-10-025-380-853/c
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429 AATTGTGCCCTTGAGGAGCTGCGTCTTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 488
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100.0%; Score 252; DB 10; Length 1915;
Best Local Similarity 100.0%; Pred. No. 2.4e-64;
Matches 252; 'Conservative 0; Mismatches 0; Indels 0;
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241 TGTGGCGCAGGA 252
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COCGANISM: Homo sapiens
US-09-964-824A-563
                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                  US-09-964-824A-101
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                                                                                                                                                                                                                                                                                           SEQ ID NO 101
LENGTH: 1915
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                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MS1-03-14
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
FRIOR FILING DATE: 2001-03-14
FRIOR FILING DATE: 2001-03-14
FRIOR PRICATION NUMBER: 60/276,026
FRIOR PRICATION NUMBER: 60/276,026
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
FRIOR PRICATION NUMBER: 60/311,732
FRIOR PRILING DATE: 2001-09-10
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-26
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FRIOR FILING DATE: 2001-09-26
FRIOR FILING DATE: 2001-09-19
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100.0%; Score 252; DB 15; Length 1907;
Best Local Similarity 100.0%; Pred. No. 2.4e-64;
Matches 252; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                       Gordon B. MILLS
Robert C. BAST, Jr.
        Shubhangi KAMATKAR
                                                                                                                                                                                                                                                             Rosemarie SCHMANDT
                                                       Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           645 rerececación 656
                                 KOVATS
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                                                                                                            Peter OLANDI
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US-10-097-340-74
                                                                                                                                                              Peter VEIBY
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Karen GLATT
                                                                                                                                                                                                                                         Karen LU
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US-09-964-824A-101
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LENGTH: 1907
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APPLICANT:
APPLICANT:
APPLICANT:
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Sequence 563, Application US/09964824A

Sequence 563, Application US/09964824A

Parent No. US20020102531A1

GENERAL INFORMATION:

ACTILE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signation TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

FILE REPERENCE: 689290-73

CURRENT APPLICATION NUMBER: US/09/964,824A

CURRENT APPLICATION NUMBER: US/60/236,033

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR PELING DATE: 2000-09-28

PRIOR PELING DATE: 2000-09-28

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Sequence 101, Application US/0964824A
 Patent No. US20020102531A1
 GENERAL INFORMATION:
 APPLICANT: Horrigan, Stephen
 TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatulation Pinvention: Sets

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Sequence 192, Application US/09967768A
Patent No. US20220150877A1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Fang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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US-09-967-768A-192
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  429 AATIGIGCCCTIGAGGAGCTGCGTCTGGGTCTTTGGGCCTCTGGGGGACCAACTCCAIGCC 488
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Patent No. US20020142981A1
Patent No. US20020142981A1
APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Unc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer FILE REFERENCE: 4921-5028-W0.
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR PILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR SEQ ID NOS: 3950
SOFTWARE PATENTIN UVET: 2.1
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US-09-880-107-3420
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US-09-967-768A-192
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LENGTH: 1915
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CURRENT APPLICATION NUMBER: US/60/236,109

PRIOR FILING DATE: 2000-09-28

PRIOR PLING DATE: 2000-09-28

PRIOR PLING DATE: 2000-09-28

PRIOR PLING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

SOFTWARE: PatentIn version 3.0

SEQ ID NO 192

LENGTH: 1915
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Patent No. US20020076414A1
GENERAL INFORMATION:
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Query Match
Best Local Similarity 100.0
Matches 252; Conservative
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ORGANISM: Homo sapiens
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                                                                                           100.0%; Score 252; DB 9; Length 1917; 100.0%; Pred. No. 2.4e-64; tive 0; Mismatches 0; Indels 0
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
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Publication No. US20020182191A1
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Macher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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Smith, Carole L.
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US-10-025-380-1105
                TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                Best Local Similarity
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                                                      US-09-922-217-1105
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LENGTH: 1917
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Birse et al.

TITLE OF INVERDATION US/10264049

PUBLICANT: Birse et al.

TITLE OF INVENTION: NUCleic Acids, Proteins, and Antibodies

FILE REFERENCE: PAIS 2002-10-04

CURRENT FILING DATE: 2002-10-04

PRIOR APPLICATION NUMBER: US/10/264,049

PRIOR FILING DATE: 2001-06-07

PRIOR APPLICATION NUMBER: US 60/209,467

PRIOR APPLICATION NUMBER: US 60/209,467

PRIOR APPLICATION NUMBER: US 60/209,467

PRIOR APPLICATION NUMBER: US 60/209,467

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SOFFWARE: PATENT OF 1360

SOFFWARE: PATENT OF 1360
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US-09-925-301-207
US-09-925-301-207
; Sequence 207, Application US/09925301
; Fatent No. US20020052308A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TILE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REPREBENCE: PA106
; CURRENT APPLICATION NURBER: US/09/925,301
; CURRENT FILING DATE: 2001-08-10
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                                                                                                                                                                                                                                                                                        0; Indels 0; Gaps
                                                                                                                                                                                                                                               Query Match

100.0%; Score 252; DB 9; Length 1996;

Best Local Similarity 100.0%; Pred. No. 2.4e-64;

Matches 252; Conservative 0; Mismatches 0; Indels 0.
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Job time : 89.8409 secs
PRIOR APPLICATION NUMBER: PCT/USOO/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 207
LENGTH: 1996
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CORGANISM: Homo sapiens
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APPLICANT: Gordon B. MILLS
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Raren LU
APPLICANT: Karen LU
APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
APPLICANT: Karen GLATT
TILLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030
                                                                                                                                                                                                                                                                                                                           Sequence 207, Apple Sequence 64, Apple Sequence 63, Apple Sequence 914, Apple Sequence 914, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 282, Apple Sequence 216, Apple Sequence 216, Apple Sequence 310, Apple Sequence 310, Apple Sequence 310, Apple Sequence 315, Apple Sequence 315, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1953, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1611, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 1953, Apple Sequence 19
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Sebastian HOBESCH
Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MCRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: John MONAHAN
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-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SOORE=pct -THR MAXEN=0
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-LONGLOG -DEV TIMEOÜT=120 -WARN TIMEOUT=30 -THRRADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGĀPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                February 12, 2004, 21:34:51 ; Search time 585.395 Seconds (without alignments) 2334.540 Million cell updates/sec
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1 MAATCEISNIFSNYFSAMYS.......YKFGKNSSGWKEEEVLQSRN 371
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Sequence 101, Application Stephen

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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat-
TITLE OF INVENTION: Sets

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FILE OF INVENTION: Sets

FRICK PILING DATE: 2000-09-28

FRICK APPLICATION NUMBER: US/60/236,028

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           996 GGCGICTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCAACTATGGGGCCAAAAGAAAAAG 1055
                                                              201 ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                281 AspileLeulleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     936 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                876 CTCGAGGGCAAGAAGAAGCACGCGCCCAGAGGCACCCACTGTGGGAGTTCATCGG
                                                                                                                                                                                                                                221 ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro
                                                                                                                                                                                                                                                                                                            756 cccacrdarddcaagcrcrrccccagcgardgrrrrcgrgacrdcaagaagggarccc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96 ATGGCTGCAACCTGTGAGATTAGCAACATTTTTTAGCAACTACTTCAGTGCGATGTACAGC 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            156 TCGGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCGATGACTTG 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81 GlutysAsntysTyrAspAlaSerAlaIleAspPheSerArgCysAspMetAspGlyAla 100
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Mismatches:
Indels:
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Matches:
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR PLILING DATE: 2002-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-03-14
PRIOR PLILOR DATE: 2001-03-14
PRIOR PLILOR DATE: 2001-03-14
PRIOR PLILOR DATE: 2001-03-14
PRIOR PLILOR DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-16
PRIOR PLING DATE: 2001-09-26
PRIOR PLING DATE: 2001-09-26
PRIOR PLING DATE: 2001-09-19
PRIOR PLING DATE: 2001-09-19
NUMBER OF SEC ID NOS: 363
SCOTTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 74
LENGTH: 1907
TYPE: DNA
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	480 CTCCHICLERICAGGACTCTCCAGCTTCTTTTTTTTTTTTTTTTTTTTTT	181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrAlaGly 200	221 ProthraspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro 240	261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheileArg 280	301 GlyvalPheLysPheLeuArgSerGluAlavalAlaglnLeuTrpGlyGlnLysLysLys 320	IleleuGlukrgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGly 360 ATCCTGGAACGGATGGATGGTTGTTTTTTTTTTTTTTTT
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CTCCAGAGTCGGAAC 1232
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Page 4

DB: US-08-978-217-2 (1-371) x US-09-880-107-3420 (1-1915) QY 1 MetalaalaThrCysGluIleSerAsnIlePheSerAsnTyrPheSerAlaMetTyrSer 20	Db 120 ATGGCTGCAACTTGTGAACATTTTTAGCAACTTCAGTGCGATGTACAGC 179 Qy 21 SerGluAspSerThrLeuAlaSerValProProAlaAlaThrPheGlyAlaAspAspLeu 40	Qy 41 ValleuThrLeuSerAsnProGlnMetSerLeuGluGlyThrGluLysAlaSerTrpLeu 60	Oy 61 GlyGluGlnProGlnPheTrpSerLysThrGlnValLeuAspTrpIleSerTyrGlnVal 80	Qy 81 GiulysasniysTyraspalaseralaileaspPheSerargCysaspWetaspGlyala 100	Oy 101 ThrLeuCysAsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGln 120	OY 121 LeudisAlaGlnLeudrgAspLeuThrSerSerSerSsrAspGluLeuSerTrpllelle 140	Oy 141 GluLeuLeuGluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAsp 160	Oy 161 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180	Qy 181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrAlaGly 200	Qy 201 ThrGlyAlaSerArgSerSerHisSerSerHisSerGerGlyGlySerAspValAspLeuAsp 220	Qy 221 ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysLysLysByro 240	Qy 241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 260	Qy 261 LeugludlyLysExserLysBisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280 [Cy 281 AspileLeuileHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300	Oy 301 GlyvalPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys 320	321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 340	Db 1080 AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCCATGAGGTACTACTACAAACGGGAG 1139 Ov 341 IleLeuGluArqValAspGlvArqArqLeuValTyrIy9FheGlyLy8AsnSerSerGly 360
Oy 181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrhlaGly 200	Db 720 ACTGGTGCTTCTCGGAGCTCCCACACTCGGTGGAAGTGACGTGGACCTGGAT 779 Qy 221 ProThrAspGlybyEcuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro 240	Qy 241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 260	Cy 261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArg 280	Qy 281 AspileLeuileHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300	Oy 301 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysIysLys 320	Qy 321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 340	Cy 341 IleLeuGluargValAspGlyArgArgLeuValTyrLysPheGlyLsAshSerSerGly 360	Cy 361 TrpLysGluGluGalleuGlnSerArgAsn 371 Db 1200 TGGAAGGAGGAAGAGTTCTCCAGAGTCGGAAC 1232	RESULT 4 US-09-880-107-3420 ; Sequence 3420, Application US/09880107	<pre>/ Patent No. USZUUZO142981A1 / GENERAL INFORMATION: / APPLICANT: Horne, Darci T. / APPLICANT: Vockley, Joseph G.</pre>	<pre>// APPLICANT: Scherf, Uwe // APPLICANT: Gene Logic, Inc. // TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer // FILE REFERENCE: 44921-5028-WO</pre>	CURRENT APPLICATION NUMBER: US/09/880,107 CURRENT FILING DATE: 2001-06-14 PRIOR APPLICATION NUMBER: US 60/211,379 PRIOR FILING DATE: 2000-06-14	; PRIOR APPLICATION NUMBER: US 60/237,054 ; PRIOR FILING DATE: 2000-10-02 ; NUMBER OF SEQ ID NOS: 3950 ; SOFTWARE: PatentIn Ver. 2.1	; SEQ ID NO 3420 ; LENGTH: 1915 ; TYPE: DNA ; ORGANISM: Homo sapiens	; FEATURE: ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843 US-09-880-107-3420	8.69e-212 Length: 1980.00 Matches:	Percent Similarity: 100.00% Conservative: 0 Best Local Similarity: 100.00% Mismatches: 0 Query Match: 100.00% Indels: 0

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960 gacarccrcarccacccaagcrcaacgaggccrcargaagrggagaarcggcargaa 1019
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                                                161 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180
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Fatera No. US2020076414A1

GENERAL INFORMATION:
APPLICANT: Ru, Jiangchun
APPLICANT: Bocrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Adeleine Joy
APPLICANT: Weagher, Madeleine Joy
APPLICANT: Wang, Tongrong
APPLICANT: Wang, Tongrong
APPLICANT: Wing, Tongrong
APPLICANT: Spick, John A.
APPLICANT: Sinik, John A.
APPLICANT: Sinik, John A.
APPLICANT: Sinik, Carole Lynn
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APPLICANT: Wing, Gordon E.
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Fatent No. US20020150877A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT AUGUSTUR.

TITLE OF INVENTION:

TITLE OF INVENTION:

CURRENT APPLICATION NUMBER: US/09/967,768A

CURRENT APPLICATION NUMBER: US/60/236,109

FRIOR FILING DATE: 2000-09-28

FRIOR FILING DATE: 2000-09-28

FRIOR APPLICATION NUMBER: US/60/236,111

FRIOR APPLICATION NUMBER: US/60/236,111

FRIOR FILING DATE: 2000-09-28

FRIOR FILING DATE: 2000-09-28

FRIOR FILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

SOFTWARE: PatentIn version 3.0

SEQ ID NO 192

LENGTH: 1915

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1140 AICCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1199
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	SULT 7 SULT 7 SQUENCE 105 ADDICATION OF 10025380 Publication No. US20020182191A1 GENERAL INFORMATION: APPLICANT: Xu, Jiangchun APPLICANT: Secrist, Heather APPLICANT: Benson, Darin R. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Mang, Tongtong APPLICANT: Mang, Tongtong APPLICANT: Stolk, John A. APPLICANT: Sich, Carole I. APPLICANT: Sich, Carole I. APPLICANT: King, Gordon E. APPLICANT: King, Gordon E. APPLICANT: King, Gordon E. APPLICANT: King, Gordon E. APPLICANT: King, Gordon E. APPLICANT: Glapper, Jonathan D. APPLICANT: Glapper, Jonathan D. APPLICANT: Gary K. APPLICANT: Gary K. APPLICANT: Gary K. APPLICANT: Gary K. APPLICANT: Gary K. APPLICANT: Mang, Ma	### TINGER S. Dartick COMPOUNDS FOR IMMUNOTHERAPY AND COMPOUNDS CANCER AND METHODS FO 1121-171(14) N NUMBER: US/10/025,380 E: 1020 S: 1129 For Windows Version 4.0 #### B: 71e-212 #### B: 71e-212 #### B: 71e-212 #### Conservative: 1080.00 #### Conservative:	Best Local Similarity: 100.00\$ Mismatches: 0
Alignment Scores: 8 .71e-212 Length: 1917 Pred. No.: 1980.00 Matches: 371 Score: 1980.00 Mismatches: 371 Best Local Similarity: 100.00\$ Mismatches: 0 Guery Match: 100.00\$ Indels: 0 Best Local Similarity: 100.00\$ Mismatches: 0 Guery Match: 9 100.00\$ Indels: 0 Best Local Similarity: 100.00\$ Mismatches: 0 Cuery Match: 9 100.00\$ Indels: 0 Cuery Match: 9 Cuery Match: 100.00\$ Cuery Mismatches: 0 Cuery Match: 100.00\$ Mismatches: 0 Cuery Mismatches: 0 Cuery Match: 100.00\$ Mismatches: 0 Cuery Mismatches: 0 Cuery Mismatches: 0 Cuery Mismatches:	Db 122 ATGGCTGCAACTGTGAATTTTTTGCAACTACTCAGGGGATGTACGC 181 21 SerGluaspSerThrLeualaSerValProProAlaalaThrPheGlyAlaaspAspLeu 40	121 LeudisAlaGlnLeuArgAspleuThrSerSerSerSerAspGluleuSerTrpIleIle 121 LeudisAlaGlnLeuArgAspleuThrSerSerSerAspGluleuSerTrpIleIle 182 CTCCATGCCGAGGAGACTTCCAGTTCTGATGAGGTCAGTTGGATGATGATGATGAGTTGATGATGATGATGATGAT	22 ACTGGTGCTTCTCGAGGCTCCCACTCCTCAGACTCCGGTGGAGGTGGCGTGGAGCTGGAT 22 ACTGGTGCTTCTCGAGGCTCCCTCAGACTCCGGTGGAGGTGGCGTGGACCTGGAT 23 DrothrapGdlyLysleuPheProSerAspGlyPheArgAspCysLyslysGlyAspPro 24 LysHisGlyLysArgLysArgGlyArgProArgLysleuSerLysGluGlyAspPro 25 LeuGluGlyLysLysCargGaAACGAGGCCCGCGGAAACTGCAAAAAGGGGGATCCC 26 LeuGluGlyLysLysSerLysHisAlaProArglyThrHisLeuTrpGluPheIleArg 26 LeuGluGlyLysLysSerLysHisAlaProArglyThrHisLeuTrpGluPheIleArg 27 CTCGAGGCAAGAAGAGCAGCCCCCGAAAGCTGACCCTGTGGGACTTGTCGG 28 AspIleLeuIleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 29 AcatccTCATCCACCCGGAGCTCAACGAGGCCCCATGAGAATCGGCATGAA 30 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysBysLys

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Publication No. US20040005579A1
GENERAL INFORMATION:
APPLICANT: Birse et al.
ITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REPERENCE: PA133P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
FRIOR APPLICATION NUMBER: PCT/US01/18569
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                  60/209,467
PRIOR FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: US 60,
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
SEQ ID NO 756
LENGTH: 1956
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201 ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp 220
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Sequence 64, Application US/2030235915A1
GENERAL INFORMATION:
APPLICANT: SPECHY, THOMAS
APPLICANT: HINZMANN, BERND
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: ROSENTHAL, ANDRE
APPLICANT: ROSENTHAL, ANDRE
APPLICANT: TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMAN APPLICANT:
TITLE OF INVENTION: TUMAN APPLICANT:
FILE REFERENCE: SCH-1763
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    261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280
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| Sequence 207, Application US/09925301
| Patent No. US20020052308A1
| GENERAL INFORMATION:
| APPLICANT: Rosen et al. |
| TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies FILE REPREENCE: PA.06
| CURRENT APPLICATION NUMBER: US/09/925,301
| CURRENT FILING DATE: 2001-08-10
| PRIOR APPLICATION NUMBER: 60/124,270
| PRIOR APPLICATION NUMBER: 60/124,270
| PRIOR FILING DATE: 1999-03-12
| NUMBER OF SEQ ID NOS: 1694
| SEQ ID NOS: 1694
| SEQ ID NO 2076
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Best Local Similarity:
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US-09-925-301-207
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US-09-922-217-853
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Best Local Similarity:
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US-09-922-217-853/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         254 GAGCTCAGTTGGATCATTGAGCTGCTGGAAAGATGGCATGGCCTTCCAGGAGGCCCTA 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AspProGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGln 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          314 GACCCAGGGCCCTTTGACCAGGGAGCCCCTTTGCCCAGGAGCTGCTGGACGACGACGGCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       493 AGTGACGTGGACCTGGATCCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGAC 552
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Matches:
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Mismatches:
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CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
FRIOR APPLICATION NUMBER: 09/646,673
FRIOR FILING DATE: 2000-09-20
FRIOR APPLICATION WUMBER: PCT/DE99/00908
FRIOR FILING DATE: 1996-03-19
NUMBER OF SEQ ID NOS: 202
SOFTWARE: Petentin Ver: 2.1
SOFTWARE: 2269
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ORGANISM: Homo sapiens
US-10-131-410-64
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Best Local Similarity:
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APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrifst, Heather
APPLICANT: Berist, Heather
APPLICANT: Berist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stilk, John A.
APPLICANT: Stilk, John A.
APPLICANT: Sinkh, Carole Lynn
APPLICANT: Sinkh, Carole Lynn
APPLICANT: Sinkh, Carole Lynn
APPLICANT: Mang, Aijun
APPLICANT: Mang, Aijun
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APPLICANT: Ting, Gordon E.
APPLICANT: Ting, Gordon E.
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US-08-978-217-2 (1-371) x US-10-025-380-853 (1-626)
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384 TCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAG 325
                                       153 AlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAsp 172
                                                                                 324 GCCCTAGACCCAGGGCCCTTTGACCAGGGCAGCCCCTTTGCCCAGGAGCTGCTGGACGAC 265
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APPLICANT: Glapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
SOFTWARE OF SEQ ID MOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
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US-09-833-263-853
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133 SerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetAlaPheGlnGlu 152
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210.121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
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SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Macaber, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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624 GGTACAGAGAGACCAGCTGGGTTGGGGGAACAGCCCCCAGTTCTGGTCGAAGACGCAGGTT 565
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53 GlyThrGluLysAlaSerTrpLeuGlyGluGlnProGlnPheTrpSerLysThrGlnVal
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APPLICANT: Xu, Janaghun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Branch, John A.
APPLICANT: Mang, Yuqiu
APPLICANT: Wang, Tongtong
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Ming, Addon E.
APPLICANT: Ming, Addon E.
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APPLICANT: Ming, Ming, Addon E.
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SEQ ID NO 944
LENGTH: 563
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Patent No. US20020076414A1
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ORGANISM: Homo sapiens
US-09-922-217-944
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Sequence 944 Application US/09833263
; Sequence 944 Application US/09833263
; Patent No. US20020110547A1
; GENERAL INFORMATION:
    APPLICANT: Glapper, Jonathan D.
    APPLICANT: Glapper, Jonathan D.
    APPLICANT: Stolk, John A.
    APPLICANT: GLOBER, John A.
    APPLICANT: Madelen J.
    APPLICANT: Madelen J.
    TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
    TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
    TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
    TITLE OF INVENTION NUMBER: US/99/833,263
    CURRENT APPLICATION NUMBER: US/99/833,263
    CURRENT FILING DATE: 2001-04-10
    NUMBER OF SEQ ID NOS: 1093
    SEOTWARE: FRASESQ for Windows Version 3.0
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Matches:
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CRGANISM: Homo sapien
US-09-833-263-944
                                                         Percent Similarity:
Best Local Similarity:
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February 13, 2004, 01:27:22; Search time 393.438 Seconds (without alignments) 10448.744 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
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SUMMARIES

Description Sequence	Seguenc	Sequence	Sequence 3	Sequence 19	Sequence 110	Sequence 11	Sequence 207	Seguence 75(Sequence 64,	Sequence 853,	Seguence 853, 7	Sequence 853	Sequence 944,	Seguence 944
ID 	US-09-964-824A-101	US-09-964-824A-563	US-09-880-107-3420	US-09-967-768A-192	US-09-922-217-1105	US-10-025-380-1105	US-09-925-301-207	US-10-264-049-756	US-10-131-410-64	US-09-922-217-853	US-09-833-263-853	US-10-025-380-853	US-09-922-217-944	US-09-833-263-944
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% Query Match Length DB	1915	1915	1915	1915	1917	1917	1996	1956	2269	626	626	626	563	563
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Sequence 944, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 2216, App Sequence 1953, App Sequence 11873, App Sequence 44, App1 Sequence 309, App	
US-10-025-380-944 US-09-604-287A-282 US-09-604-287A-282 US-09-604-287A-282 US-10-114-805-282 US-11-124-805-282 US-11-076-622-282 US-09-998-598-2890 US-09-998-598-2890 US-09-998-598-1290 US-09-98-598-1290 US-09-98-598-1290 US-09-98-598-1490 US-09-780-669-44 US-09-780-669-44 US-09-780-669-44 US-09-780-669-44 US-09-895-118-44 US-09-895-118-44 US-09-895-118-44 US-09-895-118-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44 US-09-895-814-44	
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RESULT 1

US-10-097-340-74

Sequence 74 Application US/10097340

Publication No UB20030087250A1

SEQUENCE 74 Application US/10097340

Publication No UB20030087250A1

SEGREBAL INFORMATION:

APPLICANT: Wanjula GANNAVRAPU

APPLICANT: Sebatian HORRSCH

APPLICANT: State of WEYERS

APPLICANT: Rachel E. MEYERS

APPLICANT: Machael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Michael MORRISSY

APPLICANT: Mobert C. BANT

APPLICANT: Rachel U BAST, Jr.

APPLICANT: Rachel GLIT

APPLICANT: Namel ZHAO

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APPLICANT: Namel ZHAO

APPLICANT: Numbers: US/10/097,340

CURRENT PALLON NUMBER: US/10/097,340

CURRENT PALLON NUMBER: US/226,025

PRIOR PLICATION NUMBER: US/226,025

PRIOR PLICATION NUMBER: US/226,025

PRIOR PLICATION NUMBER: US/226,026

PRIOR APPLICATION NUMBER: US/226,026

PRIOR APPLICATION NUMBER: US/226,026

PRIOR PLICATION NUMBER: US/226,026
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361 CICCATGCCCAGCTGCGAGACCTCACCTTCTGATGAGCTCAGTTGGATCATT 420
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Sequence 101, Application US/09964824A
Ferent No. US20020102531A1
GENERAL INFORMATION:
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                                                                                                                                                                                                 100.0%; Score 1116; DB 15; Length 1907; 100.0%; Pred. No. 2e-313; Affine O; Mismatches O; Indels O;
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SEQ ID NO 74
LENGTH: 1907
                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 1116; Conservative
                                                                                                                    ; TYPE: DNA; ORGANISM: Homo sapiens US-10-097-340-74
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat TITLE OF INVENTION: Sets STILE REFERENCE: 669290-73
CURRENT APPLICATION NUMBER: US/09/24,824A
CURRENT PILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
OSOFTWARE: Patentin Version 3.0
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901 GGCGTCTTCAAGTTCCTGCGCTCCCGAGGCTGTGGCCCCAACTATGGGGCCCAAAAGAAAAAG
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CURRENT APPLICATION NUMBER: US/69/236,033
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US-09-880-107-3420
                                                                                                                                                                                                                                                                                                                          APPLICANT: Horne, Darci T.
APPLICANT: Wockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REPRENCE: 44921-9028-WO
CURRENT PILIGATION NUMBER: US/99/80,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
                                                                                                                        1081 TGGAAGGAAGAAGATTCTCCAGAGTCGGAACTGA 1116
                                                                                                                                                    1200 TGGAAGGAGGAAGGTTCTCCCAGAGTCGGAACTGA 1235
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Patent No. US20020142981A1
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3420
LENGTH: 1915
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ORGANISM: Homo sapiens
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Sequence 192, Application US/09967768A

Patent No. US2002015887A1

GENERAL INFORMATION:

APPLICANT: Augustus, Meena

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat

TITLE OF INVENTION: Sets

FILE REFERENCE: 689290-72

CURRENT APPLICATION NUMBER: US/09/967,768A

CURRENT FILING DATE: 2001-09-28

PRIOR APPLICATION NUMBER: US/60/236,109

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

SOFTWARE: Patentin Version 3.0

SEQ ID NO 192

LENGTH: 1915

TYPE: DNA

GRAANISM: Home sapiens

US-09-967-768A-192
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Score 1116; DB 10; Length 1915; Pred. No. 2e-313;

100.0%;

OY 1081 TGGAAGGAAGAAGGTTCTCCAGAGTCGGAACTGA 1116 	RESULT 6 US-09-922-217-1105 ; Sequence 1105, Application US/09922217 ; Detent No. 125002007481431	GENERAL INFORMATION: APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J.	APPLICANT: Secrist, Hearner APPLICANT: Benson, Darin R. APPLICANT: Meagher, Madeleine Joy APPLICANT: Stolk, John A.) APPLICANT: Wang, Tongroup) APPLICANT: Smith, Carole Lynn) APPLICANT: King, Gordon E.	## APPLICANT: Wang, Aljun APPLICANT: Clapper, Jonathan D. TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIA TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR TH	US/09/922, US-03		in Homo sapiens 217-1105 cc	o; Mismatche TTAGCAACATTT	Db 122 ATGCTGCACCTGTGAGTTAGCACTTTTAGCACTTATTAGCACTTTTTAGCACTTGTTTTAGCACCTGCTAGCTTGCACCTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTAGCTTGCTGCTAGCTTGCTAGCTTGCTAGCTTGCTAGCTTGCTAGCTTGCTAGCTA	182	Db 242 GTACTGAGCACCCCCAGATGCATTGGAGGGGTACAGA	302	362	422		DB 482 CICCARGCCCAGCIACIACITCCAGLICLICIGALGA Qy 421 GAGCIGCTGCAGAAGGAIGGCAIGGCAIGAGCTICCAGGAGAGGCCTIAGA	Db 542 GAGCTGCTGGAGAAGGATGGCATGCCTTCCAGGAGGCCTTCTAGAGGAGGCCCTTTGCCCAGGAGGCTGCTGAGAGGAGGCTGAGAGGAGGCTGAGAGGAGGCTGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG
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Matches 1116; Conservative 0; Mismatches 0; Indels 0; Gaps 1 ATGGCTGCAACCTGTGAGTAACAACATTTTAGCAACTACTTCAGTGGGATGTACAGG 6	61 TCGGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCGATGACTTG 1	121 GTACTGACCTGAGCAACCCCCAGATGTCATTGGAGGGTACAGAGAAGGCCAGCTGGTTG 1	181 GGGGAACAGCCCCAGTTCTGGTCGAAGACGCAGGTTCTGGACTGGATCAGCTACCAAGTG 2	241 GAGAAGAACAAGTACGAAGCGCCATTGACTTCTCACGATGTGACATGGGGGC 3	301 ACCCTCTGCAATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGACCAA 3	361 CTCCATGCCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATT 4	421 GAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGAC 4	481 CAGGGCAGCCCTTTGCCCAGGAGCTGCTGGACGGCGGCAGGCA	541 CCCGGCAGCTGTGGCGCAGGAGCCCCCTCCCCTGGCAGCTCTGACGTCTCCACCGCAGGG (6	601 ACTGGTGCTTCTCGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTGGAT (CCCACTGATGGCAAGGTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGGGGGATCCC		TICGAGGGCAAGAGGGAAGCGGCCCCCGGAAGCCTGAGCGAAGAGGTTCAGGAGCTTCTCGGCAGCCAAGAGCAAGCCAAGCCAAGCCAAGCCAAGCCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAACCAACAA	GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA	GECGICTICALCCACCCGGAGCICAGGGGCCCAACGAGGGCCCAACGAGGAGAAGAAAGA	GGCGICTICAMGITCTTGCGCTCCGAGGCTGTGGCCCAACTATGGGGCCAAAAAAA	961 AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG	1021 AICCIGGAACGGGGGAIGGCCGGCGACTCGICTACAAGITIGGCAAAAACTCAAGGGGCCCCCCCCCC
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TYPE: DNA
CRGANISM: Homo sapiens
US-10-025-380-1105
  LENGTH: 1917
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APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Scolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Mang, Yugiu
APPLICANT: Mith, Carole L.
APPLICANT: Mith, Carole L.
APPLICANT: Mith, Carole L.
APPLICANT: Gridy, Aljun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Carter, Darines S.
APPLICANT: Carter, Darines S.
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APPLICANT: Carter, Darines S.
APPLICANT: Vedvick Thomas S.
APPLICANT: Carter, Darines S.
APPLICANT: VENNENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOSIS
TITLE OF INVENTION: OLD COLON CANDER AND METHODS FOR THEIR USE
CURRENT APPLICATION NUMBER: US/10/025, 380
CURRENT APPLICATION NUMBER: US/10/12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
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Publication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: Lodes, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Wangher, Madeleine Joy
APPLICANT: Wang, Tongtong
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sitlk, Cardole I.
APPLICANT: Smith, Cardole I.
APPLICANT: King, Gordon E.
APPLICANT: Mang, Aljunian
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Query Match 100.0%; Score 1116; DB 14; Length 1917; Best Local Similarity 100.0%; Pred. No. 2e-313; Matches 1116; Conservative 0; Mismatches 0; Indels 0;
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Db 621 CAGGGCAGCTTTTGCCCAGGAGCTGCTGGAAGCCAAGCC	221 AAGCACGGAAACGACCACCCGAAAGCTGTGCCAAAGAAAG	Qy 1021 ATCTGGAACGGTGGATGGCCGCGACTCGTCTACAAAATTTGGCAAAAACTCAAGGGGC 1080	RESULT 9 US-10-264-049-756 US-10-264-049-756 Sequence 756, Application US/10264049 Publication No. US20040005579A1 GENERAL INFORMATION: APPLICAMIN: Bitse et al. TITLE OF INVENTION: CURRENT ELING DATE: 2002-10-04 PRIOR PILING DATE: 2002-10-04 PRIOR PELING DATE: 2000-06-07 PRIOR PELING DATE: 2000-06-07 PRIOR PILING DATE: 200-06-07 PRIOR PILING DATE: 2000-06-07 PRIOR PILING DATE: 2000-06-0
Qy 961 AACAGCAACATGACCTACGAGAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 1020 Db 1082 AACAGCAACATGACCTACGAGAAGCTGAGCCGTGAGGTACTACTACAAACGGGAG 1141 CD 1021 ATCCTGGAACTGAGCTGAGCTGGTCGTCTACAAGTTTGGCAAAAACTCAAGGGGC 1080 Db 1142 ATCCTGGAACGGGTGGATGGCGGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1201 CO 1111	RESULT 8 US-09-925-301-207 Squence 207, Application US/09925301 Fatent No. US2002052308A1 Fatent No. US2002052308A1 GENERAL INFORMATION: TITLE OF INVENTION: TITLE OF INVENTION NUCLEIC Acids, Proteins and Antibodies CURRENT PILING DATE: 2001-08-10 FRIOR PELICATION NUMBER: US/09/925,301 CURRENT FILING DATE: 2000-03-08 FRIOR FILING DATE: 2000-03-08 FRIOR FILING DATE: 2000-03-08 FRIOR FILING DATE: 1999-03-12 NUMBER OF SEQ ID NOS: 1694 SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 207 LENGTH: 1996 TYPE: DNA TYPE: DNA TYPE: DNA TYPE: DNA SOFTWARE: NA SOFTWARE	Query Match Best Local Similarity 99.9%; Pred. No. 2.7e-313; Matches 1115; Conservative 1; Mismatches 0; Indels 0; Gaps 0; Qy 1 ATGGCTGCAACCTGTGAGATTAGCAACATTTTAGCAACTACTTCAGTGCGATGTACAC 60 Db 141 ATGGCTGCAACCTGTGAGATTAGCAACATTTTAGCAACTACTTCAGTGCGATGTACAC 60	0

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ACTCCGGTGGAAGTGACGTGGACCTGGATCCCACTGATGGCAAGCTCTTCCCCAGCGATG 691
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                                                                                                                                                                                                                     APPLICANT: SPECHT, THOMAS
APPLICANT: SPECHT, THOMAS
APPLICANT: SPECHT, THOMAS
APPLICANT: HINZMANN, BERND
APPLICANT: HINZMANN, BERND
APPLICANT: PILARSKY, CHARISTIAN
APPLICANT: PILARSKY, CHARISTIAN
APPLICANT: POGAR
APPLICANT: POGAR
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAD
TITLE OF INVENTION: HUMARS
TITLE OF INVENTION: HUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
PRIOR PILING DATE: 2000-09-20
PRIOR PILING DATE: 1099-00-19
PRIOR FILING DATE: 1999-00-19
PRIOR FILING DATE: 1999-00-19
SEOFTH PRIOR FILING DATE: 1999-00-19
SEOFTH PRIOR FILING DATE: 1999-00-19
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SEOFTH P
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                               US-10-131-410-64
Sequence 64, Application US/10131410
Publication No. US20030235915A1
GENERAL INFORMATION:
APPLICANT: SPECHT, THOMAS
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; FILE REFERENCE: 210121.471C13
; CURRENT APPLICATION NUMBER: US/09/922,217
; CURRENT FILING DATE: 2001-08-03
; NUMBER OF SEG ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
TYPE: N."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 853, Application US/09922217
Patent No. US20020076414A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lodes, Michael J.
Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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APPLICANT: Xu, Jiangchun
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ORGANISM: Homo sapiens
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APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND
TITLE OF APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT PILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
NUMBER OF SEQ ID NOS: 1093
SOPTWARE: FastSEQ for Windows Version 3.0
446 iGSICITITGGGCCICIGGGGGACCAACICCAIGCCCAGCIGCGAGACCICACIICCAGCI
                                                                                                             395 CTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGGAAAGGATGGCATGGCCTTCCAGG
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99.8%; Pred. No. 6.5e-171;
live 0; Mismatches 1;
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755 AGCIGAGCAAAGAGIACIGGGACIGI 780
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Secrist, Heather
Secrist, Heather
Benson, Darin R.
Meagher, Madleine Joy
Stolk, John A.
Wang, Tongtong
Jang, Yuqiu
Smith, Carole Lynn
King, Gordon E.
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CORGANISM: Homo sapiens
US-09-922-217-944
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                                                          395 CTICTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGG 454
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446 TGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCCCAGCTGCGAGACCTCACCTTCCAGCT 387
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
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55.9%; Score 624.4; DB 14;
Best Local Similarity 99.8%; Pred. No. 6.5e-171;
Matches 625; Conservative 0; Mismatches 1;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
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APPLICANT: Lodes, Michael J.
APPLICANT: Becrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Mang, Tongtong
APPLICANT: Ming, Yugiu
APPLICANT: Smith, Carole L.
APPLICANT: King, Gordon E.
APPLICANT: King, Aljun
APPLICANT: King, Aljun
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Vedyick Thomas S.
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ORGANISM: Homo sapiens
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TICTGGACTGGATCAGCTACCAAGTGGAGAACAAGTACGACGCAAGCGCCATTGACT 274
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